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OM protein - protein search, using sw model

Run on: February 27, 2003, 14:15:13 ; Search time 35 Seconds
(without alignments)
137.058 Million cell updates/sec

Title: US-09-634-363-2

Perfect score: 194

Sequence: 1 YPIKPEAPGEDASPELNRYASLRHYNLVTRQRY 36

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 908470 segs, 133250620 residues

Total number of hits satisfying chosen parameters: 908470

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : A_Geneseq_101002.*

- 1: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1980.DAT.*
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- 4: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1983.DAT.*
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- 13: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1992.DAT.*
- 14: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1993.DAT.*
- 15: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1994.DAT.*
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- 19: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1998.DAT.*
- 20: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1999.DAT.*
- 21: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA2000.DAT.*
- 22: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA2001.DAT.*
- 23: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA2002.DAT.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	194	100.0	36	15	Human peptide YY (
2	194	100.0	36	17	Human peptide YY.
3	194	100.0	36	19	Human peptide YY.
4	194	100.0	36	20	Peptide Y. Synthe
5	194	100.0	36	21	Human peptide YY.
6	194	100.0	36	21	Human neuropeptide
7	194	100.0	36	21	Human peptide YY (
8	194	100.0	36	21	Human peptide YY (
9	194	100.0	36	22	Peptide YY SEQ ID
10	194	100.0	36	23	Human peptide tyro
					Modified human pep

11	194	100.0	97	21	AAB08020	Amino acid sequenc
12	194	100.0	176	22	AABG75364	Human colon cancer
13	188	96.9	36	22	AAB91109	Parathyroid hormon
14	185	95.4	36	22	AAB91226	Peptide YY SEQ ID
15	184	94.8	36	15	AAB62049	Porcine peptide YY
16	184	94.8	36	17	AAR97740	Porcine peptide YY
17	184	94.8	36	18	AAW15365	[Am-DNP-His26]-PYY
18	184	94.8	36	19	AAW51800	Porcine peptide YY
19	184	94.8	36	21	AAW87549	Porcine peptide YY
20	184	94.8	36	22	AAB91225	Peptide YY SEQ ID
21	184	94.8	36	22	AAU06187	Porcine peptide ty
22	180	92.8	34	22	AAB91224	Peptide YY SEQ ID
23	177	91.2	36	11	AAR07278	Porcine small inte
24	175	90.2	36	19	AAW51808	Peptide YY analogu
25	168	86.6	36	11	AAW07277	Porcine small inte
26	166	85.6	36	20	AAW50293	Neutrophil-activat
27	160	82.5	36	11	AAW07276	Porcine small inte
28	155	79.9	36	16	AAR87890	Neuropeptide Y ago
29	155	79.9	36	22	AAB91222	Pancreatic polypep
30	146	75.3	36	17	AAW06955	Porcine neuropepti
31	146	75.3	36	17	AAW97743	Porcine neuropepti
32	146	75.3	36	21	AAW51550	Rat and porcine ne
33	146	75.3	36	22	AAB97621	Porcine/rat neurop
34	144	74.2	36	17	AAR97742	Human neuropeptide
35	144	74.2	36	19	AAW51823	Peptide YY analogu
36	144	74.2	36	21	AAW51549	Human neuropeptide
37	144	74.2	36	22	AAE06684	Human neuropeptide
38	144	74.2	36	22	AAB97620	Human neuropeptide
39	144	74.2	36	22	AAB91213	Neuropeptide Y pep
40	144	74.2	36	22	AAU06195	Mammalian neuropep
41	144	74.2	36	23	AAU85985	Modified human neu
42	144	74.2	97	20	AAW43334	Neuropeptide Y. S
43	144	74.2	97	20	AAW23828	Human prepro-neuro
44	144	74.2	97	21	AAB35660	Human neuropeptide
45	144	74.2	97	21	AAW57078	Human neuropeptide

ALIGNMENTS

RESULT 1

AAR62050
ID AAR62050 standard; peptide; 36 AA.

XX AAR62050;

XX AC

XX 14-JUN-1995 (first entry)

DT Human peptide YY (PYY).

DE Peptide YY; PYY; gastro-enterological disorders; intestinal water;

XX electolyte secretion; cell proliferation; nutrient transport;

XX lipolysis; blood flow regulation.

OS Homo sapiens.

XX WO9422467-A.

XX 13-OCT-1994.

XX 29-MAR-1994; 94WO-US03380.

XX 29-MAR-1993; 93US-0038534.

XX 19-AUG-1993; 93US-0109326.

XX (UYCI-) UNIV CINCINNATI.

XX Balasubramaniam A;

XX WPI; 1994-332815/41.

XX New peptide derivs. - useful as therapeutic agents, for treating

PT gastro-enterological disorders

XX Disclosure; Page 3; 45pp; English.

PS

CC AAR62050 describes the amino acid sequence of human peptide YY (PYY), which was isolated from the endocrine cells of the human gastrointestinal tract and pancreas. Using the equivalent porcine PYY sequence (AAR62049) as a base the PYY analogues described in CC AAR62051-R62082 were produced. The new peptides were found to have a variety of properties, that made them useful as therapeutic agents in the treatment of gastro-enterological disorders. As part of a therapeutic composition they could be used for decreasing CC excess intestinal water and electrolyte secretion, for regulating CC cell proliferation and augmenting nutrient transport, and for CC regulating lipolysis and blood flow.

XX Sequence 36 AA;

SQ

Query Match 100.0%; Score 194; DB 15; Length 36;
Best Local Similarity 100.0%; Pred. No. 2.7e-20;
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 YPIKPEAGEDASPEELNRYASLRHYNLVTRQRY 36
|||||
Db 1 YPIKPEAGEDASPEELNRYASLRHYNLVTRQRY 36

RESULT 2

AAR97741

ID AAR97741 standard; peptide; 36 AA.

XX

AC AAR97741;

XX

DT 10-JAN-1997 (first entry)

XX

DE Human peptide YY.

XX

XX Peptide YY; PYY; porcine; human; intestine; endocrine cell; gut motility; gastrointestinal tract; pancreas; inhibitor; intestinal secretion; pig; pancreatic tumour; blood flow; serous cyst adenoma; microcystic tumour; Solid-cyst tumour; malignant tumour; therapy.

XX

OS Homo sapiens.

XX

XX WO9614854-A1.

PN

XX

XX 23-MAY-1996.

PD

XX

XX 03-NOV-1995; 95WO-US14303.

PF

XX

XX 14-NOV-1994; 94US-0338395.

PR

XX

XX (REGC) UNIV CALIFORNIA.

PA

XX

PI Mcfadden DW;

XX

XX WPI; 1996-259558/26.

DR

XX

XX Use of peptide YY and its agonists to treat pancreatic tumours - either in vitro or in vivo to reduce tumour cell proliferation

PT

XX

PS Disclosure; Page 3; 22pp; English.

XX

XX AAR97740 and AAR97741 represent porcine and human peptide YY (PYY) respectively. This sequence is isolated from intestine, and is localised in the endocrine cells of the gastrointestinal tract and the pancreas. PYY is thought to inhibit gut motility and blood flow, to mediate intestinal secretion, and stimulate net absorption. These sequences, and agonists against them (see AAR97742-R97744), can be used in the method of the invention. The method of the invention is for CC inhibiting pancreatic tumours by contacting them with an effective amount of one of these sequences. The method is effective in treating both CC benign and malignant pancreatic tumours. The types of benign tumour CC pancreatic tumours that can be treated, include, serous cyst adenomas,

CC microcystic tumours, and solid-cyst tumours. The malignant tumours capable of being treated by the method of the invention include, CC carcinomas arising from the ducts, acini, or islets of the pancreas.

XX

SQ Sequence 36 AA;

Query Match 100.0%; Score 194; DB 17; Length 36;
Best Local Similarity 100.0%; Pred. No. 2.7e-20;
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 YPIKPEAGEDASPEELNRYASLRHYNLVTRQRY 36
|||||
Db 1 YPIKPEAGEDASPEELNRYASLRHYNLVTRQRY 36

RESULT 3

AAW51801

ID AAW51801 standard; peptide; 36 AA.

XX

AC AAW51801;

XX

DT 13-OCT-1998 (first entry)

XX

DE Human peptide YY.

XX

XX peptide YY; cell proliferation; nutrient transport; lipolysis; electrolyte secretion; anti-secretory; intestinal water; antimotility.

KW

KW Homo sapiens.

OS

XX

XX WO9820885-A1.

PN

XX

XX 22-MAY-1998.

PD

XX

XX 13-NOV-1996; 96WO-US18374.

PF

XX

XX 13-NOV-1996; 96WO-US18374.

PR

XX

XX (UYCI-) UNIV CINCINNATI.

PA

XX

PI Balasubramaniam A;

XX

XX WPI; 1998-322327/28.

DR

XX

XX New analogue(s) of peptide YY - used, e.g. to control cell proliferation, nutrient transport, lipolysis and intestinal water and electrolyte secretion

PT

XX

PS Disclosure; Page 3; 54pp; English.

XX

XX The invention relates to peptide YY analogues which may be used e.g. for decreasing excess intestinal water and electrolyte secretion in mammals, CC to regulate cell proliferation (especially intestinal cell proliferation), to increase nutrient transport, to regulate lipolysis CC and to regulate blood flow. The peptides exhibit antisecretory and CC antimotility properties and are especially useful in treatment of CC gastrointestinal disorders associated with excess intestinal electrolyte CC and water secretion as well as decreased absorption. The new peptides CC are truncated versions of peptide YY. They interact solely with peptide CC YY receptors and not with homologous receptors such as NPY Y1 and Y3, CC thus minimising unwanted (ant)agonist side reactions. The present CC sequence represents human peptide YY.

XX

SQ Sequence 36 AA;

Query Match 100.0%; Score 194; DB 19; Length 36;
Best Local Similarity 100.0%; Pred. No. 2.7e-20;
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 YPIKPEAGEDASPEELNRYASLRHYNLVTRQRY 36
|||||
Db 1 YPIKPEAGEDASPEELNRYASLRHYNLVTRQRY 36

```

RESULT 4
AAY43335
ID AAY43335 standard; peptide; 36 AA.
XX
AC AAY43335;
XX
DT 25-JAN-2000 (first entry)
XX
DE Peptide Y.
XX
KW Neuropeptide Y: NPY; agonist; peptide YY; PYY; airway restriction;
KW bronchial disease; asthma; bronchitis; laryngitis; Alzheimer's disease;
KW chronic rhinosinusitis; oedema; inflammation; congestive heart failure;
KW cardiomyopathy; coronary arterial disease; myocardial infarction; AIDS;
KW diminished cardiac vagal activity; hypertension; epilepsy; ischaemia;
KW angina; immune response; antihistamine; therapy.
XX
OS Synthetic.
XX
PN WO9951626-A2.
XX
PD 14-OCT-1999.
XX
PF 26-MAR-1999; 99WO-EP02076.
XX
PR 03-APR-1998; 98US-0054393.
XX
PA (BMRA-) BMRA CORP BV.
XX
PI Mutter M, Lacroix J, Grouzmann E;
XX
DR WPI; 1999-620192/53.
XX
PT New agonists of neuropeptide Y containing linear peptide linked to
PT cyclic template peptide, used e.g. to reduce airway restriction in
PT asthma.
XX
PS Disclosure; Page 42; 45pp; English.
XX
CC This sequence represents peptide Y (PPY). The invention relates to
CC neuropeptide Y (NPY) agonists comprising: (i) a template comprising a
CC cyclic peptide (Ia) of 3-10 amino acids (aa) in which at least two
CC residues are joined by a naphthyl ring; and (ii) at least one linear
CC peptide (Ib) of 12-37 aa, bound to (i). The agonists, also NPY itself,
CC the related sequence PYY and PYY agonists, are used to reduce airway
CC restriction in patients with bronchial disease, especially asthma and
CC bronchitis. The agonists may also be used: (i) to treat conditions
CC responsive to NPY or PYY, e.g. laryngitis, chronic rhinosinusitis,
CC oedema, inflammation, anxiety, congestive heart failure, cardiomyopathy,
CC coronary arterial disease, diminished cardiac vagal activity,
CC hypertension, Alzheimer's disease, epilepsy, ischaemia, angina,
CC myocardial infarction, acquired immune deficiency syndrome and diseases
CC characterised by reduced immune responses; and (ii) to increase body
CC weight or as an antihistamine. The template induces folding of (Ib) into
CC a biologically active form. Since (i) contain only the C-terminal region
CC of NPY, they are selective for the Y2 receptor, i.e. they do produce the
CC side effects associated with binding to the Y1 receptor.
XX
SQ Sequence 36 AA;
Query Match 100.0%; Score 194; DB 20; Length 36;
Best Local Similarity 100.0%; Pred. No. 2.7e-20;
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 YPIKPEAPGEDASPEELNRYASLRHYLNLVTRQRY 36
Db 1 YPIKPEAPGEDASPEELNRYASLRHYLNLVTRQRY 36
RESULT 5
AAB12178
ID AAB12178 standard; peptide; 36 AA.

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XX AAB12178;
XX
DT 20-JUN-2001 (first entry)
XX
DE Human peptide YY.
XX
KW Human; neuropeptide Y; luteinizing hormone; reproductive system;
KW NPY; NPY-Y4 receptor; precocious puberty;
KW polycystic ovary syndrome; endometriosis; benign prostatic hyperplasia;
KW delayed puberty; amenorrhea; breast cancer; prostate cancer;
KW peptide YY; PYY.
XX
OS Homo sapiens.
XX
FH Key Location/Qualifiers
FT Modified-site 36
FT /note= "C-terminal amide"
XX
PN WO200030674-A1.
XX
PD 02-JUN-2000.
XX
PF 26-NOV-1999; 99WO-GB03963.
XX
PR 26-NOV-1998; 98GB-0025969.
PR 13-MAY-1999; 99GB-0011178.
XX
PA (FEER ) FERRING BV.
XX
PI Broqua P, Akinsanya K, Hayward A;
XX
DR WPI; 2000-399931/34.
XX
PT Treating human reproductive disorders such as amenorrhea, delayed
PT puberty, polycystic ovary syndrome and endometriosis, comprises
PT administering a neuropeptide Y-Y4 receptor ligand.
XX
PS Disclosure; Page 2; 17pp; English.
XX
CC Neuropeptide Y (NPY) (AAB12177) has a number of effects on the
CC reproductive system. NPY is one of a family of neuropeptides. Other
CC members of the family include the present sequence, peptide YY (PYY), and
CC pancreatic polypeptide (PP, see AAB12179 and AAB12180). Selective NPY-Y4
CC receptor agonists have been found (see AAB12181 to AAB12183). The NPY-Y4
CC receptor agonists cause an increase in the circulating levels of
CC luteinizing hormone (LH) and hence improve the fertility of animals with
CC compromised reproductive function. The NPY-Y4 agonists may be used to
CC treat human reproductive disorders such as delayed puberty and
CC amenorrhea. In addition, NPY-Y4 antagonists may be used to treat human
CC reproductive disorders such as precocious puberty, endometriosis,
CC polycystic ovary syndrome, benign prostatic hyperplasia and
CC hormone-dependent neoplasias e.g. breast cancer and prostate cancer. The
CC present sequence was used in a sequence homology comparison.
XX
SQ Sequence 36 AA;
Query Match 100.0%; Score 194; DB 21; Length 36;
Best Local Similarity 100.0%; Pred. No. 2.7e-20;
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 YPIKPEAPGEDASPEELNRYASLRHYLNLVTRQRY 36
Db 1 YPIKPEAPGEDASPEELNRYASLRHYLNLVTRQRY 36
RESULT 6
AAY87961
ID AAY87961 standard; peptide; 36 AA.
XX
AC AAY87961;
XX
DT 18-SEP-2000 (first entry)

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XX OS Homo sapiens.
XX OS Synthetic.
XX PN WO200069900-A2.
XX XX
XX PD 23-NOV-2000.
XX XX
XX PF 17-MAY-2000; 2000WO-US13576.
XX XX
XX PR 17-MAY-1999; 99US-0134406.
XX PR 10-SEP-1999; 99US-0153406.
XX PR 15-OCT-1999; 99US-0153783.
XX XX
XX PA (CONJ-) CONJUCHEM INC.
XX PI Bridon DP, Ezrin AM, Milner PG, Holmes DL, Thibadeau K;
XX DR WPI; 2001-112059/12.
XX XX
XX PT Modifying and attaching therapeutic peptides to albumin prevents
XX PT peptidase degradation, useful for increasing length of in vivo activity
XX PT
XX PS Disclosure: Page 327-328; 733pp; English.
XX XX
XX CC The present invention describes a modified therapeutic peptide (I)
XX CC comprising a therapeutically active amino acid region (III) and a
XX CC reactive group (II) (e.g. succinimidyl and maleimido groups) attached to
XX CC a less therapeutically active amino acid region (IV), which covalently
XX CC bonds with amino/hydroxyl/thiol groups on blood components to form a
XX CC peptidase stabilised therapeutic peptide composed of 3-50 amino acids.
XX CC (I) are useful for modifying therapeutic peptides e.g. hormones, growth
XX CC factors and neurotransmitters, to protect them from peptidase activity
XX CC in vivo for the treatment of various disorders. Endogenous therapeutic
XX CC peptides are not suitable as drug candidates as they require frequent
XX CC administration due to rapid degradation by peptidases in the body.
XX CC Modifying and attaching therapeutic peptides to albumin prevents or
XX CC reduces the action of peptidases to increase length of activity (half
XX CC life) and specificity as bonding to large molecules decreases
XX CC intracellular uptake and interference with physiological processes.
XX CC AAB90829 to AAB92441 represent peptides which can be used in the
XX CC exemplification of the present invention.
XX XX
XX SQ Sequence 36 AA;
    Query Match 100.0%; Score 194; DB 22; Length 36;
    Best Local Similarity 100.0%; Pred. No. 2.7e-20;
    Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPEELNRYASLRHYLNLVTQRQY 36
    |||||
Db 1 YPIKPEAPGEDASPEELNRYASLRHYLNLVTQRQY 36
    |||||

RESULT 9
AAU06188
ID AAU06188 standard; peptide; 36 AA.
XX AC
XX AC AAU06188;
XX DT 04-DEC-2001 (first entry)
XX XX
XX DE Human peptide tyrosine-tyrosine (PYT).
XX XX
XX KW Human; brain aluminium concentration; central nervous system;
XX KW CNS; peptide tyrosine-tyrosine receptor; PYT receptor; PP receptor;
XX KW pancreatic polypeptide receptor; Alzheimer's disease; neurotropic;
XX KW neuro protective.
XX XX
XX OS Homo sapiens.
XX PN WO200158409-A2.

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XX XX 16-AUG-2001.
XX PD
XX XX
XX PF 07-FEB-2001; 2001WO-US03952.
XX XX
XX PR 08-FEB-2000; 2000US-0499980.
XX XX
XX PA (UYNC-) UNIV NORTH CAROLINA STATE.
XX XX
XX PI Croom WJ, Berg BM, Taylor IL;
XX XX
XX DR WPI; 2001-550001/61.
XX XX
XX PT Reducing aluminium levels in the central nervous system, for the
XX PT treatment of Alzheimer's disease comprises administration of a peptide
XX PT tyrosine receptor agonist or a pancreatic polypeptide receptor agonist
XX PT
XX PS Disclosure: Page 7; 52pp; English.
XX XX
XX CC The present invention relates to a method of reducing aluminium levels
XX CC in the central nervous system (CNS). The method comprises administration
XX CC of a peptide tyrosine-tyrosine (PYT) receptor agonist or a pancreatic
XX CC polypeptide (PP) receptor agonist. The method is useful for the
XX CC treatment of Alzheimer's disease and for reducing aluminium levels in
XX CC the central nervous system, especially the brain, of a subject. The
XX CC treatments are effective and do not impart excessive toxicological
XX CC effects. The present sequence represents human PYT.
XX XX
XX SQ Sequence 36 AA;
    Query Match 100.0%; Score 194; DB 22; Length 36;
    Best Local Similarity 100.0%; Pred. No. 2.7e-20;
    Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPEELNRYASLRHYLNLVTQRQY 36
    |||||
Db 1 YPIKPEAPGEDASPEELNRYASLRHYLNLVTQRQY 36
    |||||

RESULT 10
AAU85987
ID AAU85987 standard; peptide; 36 AA.
XX XX
XX AC AAU85987;
XX DT 21-MAY-2002 (first entry)
XX XX
XX DE Modified human peptide YY.
XX XX
XX KW Increased biological potency; prolonged activity; increased half-life;
XX KW glucose intolerance; insulin resistance; type II diabetes; bone disease;
XX KW cancer; inflammatory disorder; obesity; developmental disorder;
XX KW hyperproliferative skin disease; hormone-dependent disease; homeostasis;
XX KW intestinal disease; interleukin-8 production; smooth muscle contraction;
XX KW feeding; blood pressure; pancreatic secretion; mutant; mutein; human;
XX KW peptide YY.
XX XX
XX OS Homo sapiens.
XX OS Synthetic.
XX XX
XX FH Key Location/Qualifiers
XX FT Modified-site 1
XX FT /note= "H-Tyr"
XX FT Modified-site 36
XX FT /note= "C-terminal amide"
XX XX
XX PN WO200210195-A2.
XX XX
XX PD 07-FEB-2002.
XX XX
XX PF 02-AUG-2001; 2001WO-CA01119.
XX XX

```

PR 02-AUG-2000; 2000US-222619P.
 XX (THER-) THERATECHNOLOGIES INC.
 XX
 XX Gravel D, Habi A, Abribat T;
 PI
 XX WPI: 2002-206179/26.
 DR
 XX
 XX Novel modified biological peptide with increased biological potency,
 PT prolonged activity, increased half-life, for treating glucose
 PT intolerance associated or not with insulin resistance pathologies, type
 PT II diabetes
 XX
 XX Claim 5; Page 62; 77pp; English.
 PS
 XX The present invention relates to modified biological peptides with
 CC increased biological potency, prolonged activity and/or increased
 CC half-life. The peptides of the invention are useful in the treatment
 CC of glucose intolerance which may be associated with insulin resistance
 CC pathologies, and in the treatment of type II diabetes. They are also
 CC useful for treating bone diseases, cancer, diseases related to
 CC inflammatory responses, obesity, autism, pervasive developmental
 CC disorders, hyperproliferative skin diseases, hormone-dependent diseases,
 CC They can be used for regulating blood glucose, enhancing mucosal
 CC regeneration in patients with intestinal diseases, inhibition of
 CC interleukin-8 production, stimulation of acid release, homeostasis,
 CC regulation of exocrine and endocrine secretions, smooth muscle
 CC contraction, feeding, blood pressure, body temperature and cell growth,
 CC regulation of food intake and energy balance, and stimulation of
 CC pancreatic secretion or cell growth. AAU85971-AAU86019 represent the
 CC modified biological peptides of the invention.
 XX
 XX Sequence 36 AA;
 SQ
 Query Match 100.0%; Score 194; DB 23; Length 36;
 Best Local Similarity 100.0%; Pred. No. 2.7e-20;
 Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 YPIKPEAPGEDASPEELNRYASLRHYLNLVTRQRY 36
 DB 1 YPIKPEAPGEDASPEELNRYASLRHYLNLVTRQRY 36
 RESULT 11
 AAB08020
 ID AAB08020 standard; Protein; 97 AA.
 XX
 AC AAB08020;
 XX
 DT 14-NOV-2000 (first entry)
 XX
 DE Amino acid sequence of a human peptide YY (PYY).
 XX
 KW Peptide YY; PYY; pancreatic cell growth; pancreatic tissue degeneration;
 KW glucose metabolism; insulin resistance; glucose intolerance;
 KW glucose non-responsiveness; hyperglycemia; obesity; hyperlipidemia;
 KW hyperfiltration; type II diabetes mellitus.
 XX
 OS Homo sapiens.
 XX
 FH Key Location/Qualifiers
 FT Peptide 1..28
 FT Protein /note= "signal peptide"
 FT /note= "mature protein"
 XX
 XX WO200047219-A2.
 PN
 XX 17-AUG-2000.
 PD
 XX 10-FEB-2000; 2000WO-US03391.
 PF
 XX 10-FEB-1999; 99US-0119577.
 PR

XX (ONTO-) ONTOGENY INC.
 XX
 XX Pang K, Lu H;
 PI
 XX WPI: 2000-565257/52.
 DR
 XX N-PSDB; AAA59713.
 DR
 XX Promoting the growth of pancreatic cells and reducing degeneration of
 PT pancreatic tissue for treating a disease associated with altered
 PT glucose metabolism comprises contacting with a composition including
 PT (an agonist of) peptide YY
 XX
 XX Disclosure; Page 82-83; 83pp; English.
 PS
 XX The present sequence represents a human peptide YY (PYY). PYY triggers
 CC gain of function in glucose non-responsive foetal and adult islets which
 CC leads to glucose responsivity. The specification describes a method for
 CC promoting the growth of pancreatic cells and reducing degeneration of
 CC pancreatic tissue. The method comprises contacting pancreatic cells
 CC or tissue with a composition including PYY or an agonist of PYY. The
 CC method is used for treating a disease, especially in a human, associated
 CC with altered glucose metabolism, especially insulin resistance, glucose
 CC intolerance or glucose non-responsiveness, hyperglycemia, obesity,
 CC hyperlipidemia, hyperfiltration or type II diabetes mellitus.
 XX
 XX Sequence 97 AA;
 SQ
 Query Match 100.0%; Score 194; DB 21; Length 97;
 Best Local Similarity 100.0%; Pred. No. 8.9e-20;
 Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 YPIKPEAPGEDASPEELNRYASLRHYLNLVTRQRY 36
 DB 29 YPIKPEAPGEDASPEELNRYASLRHYLNLVTRQRY 64
 RESULT 12
 AAG75364
 ID AAG75364 standard; Protein; 176 AA.
 XX
 AC AAG75364;
 XX
 DT 03-SEP-2001 (first entry)
 XX
 DE Human colon cancer antigen protein SEQ ID NO:6128.
 XX
 KW Human; colon cancer; colon cancer antigen; diagnosis; detection;
 KW colorectal carcinoma; chromosome 17.
 XX
 OS Homo sapiens.
 XX
 PN WO200122920-A2.
 XX
 PD 05-APR-2001.
 XX
 PF 28-SEP-2000; 2000WO-US26524.
 XX
 XX 29-SEP-1999; 99US-0157137.
 PR 03-NOV-1999; 99US-0163280.
 XX
 XX (HUMA-) HUMAN GENOME SCI INC.
 PA
 XX Ruben SM, Barash SC, Birse CE, Rosen CA;
 PI
 XX WPI: 2001-235357/24.
 DR
 XX N-PSDB; AAH34769.
 XX
 XX Nucleic acids encoding 4277 human colon cancer-associated polypeptides,
 PT useful for preventing, diagnosing and/or treating colorectal cancers -
 XX Claim 11; Page 7579-7580; 9803pp; English.
 PS
 XX

AAH32943 to AAH37195 and AAG7788 represent human colon cancer-associated nucleic acid molecules (N) and proteins (P), where the proteins are collectively known as colon cancer antigens. The colon cancer antigens have cytostatic activity and can be used in gene therapy and vaccine production. N and P may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate P expression. For example, N and P may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of P by expressing P. Inactive proteins or to supplement the patients own production of P. Additionally, N may be used to produce the colon cancer-associated Ps, by inserting the nucleic acids into a host cell and culturing the cell to express the proteins. N and P can be used in the prevention, diagnosis and treatment of colorectal carcinomas and cancers. AAH37196 to AAH37204 and AAB7789 represent sequences used in the exemplification of the present invention.

N.B. Pages 666 to 682 and page 7053 of the sequence listing were missing at time of publication, meaning no sequences are present for SEQ ID NO:1027 to 1052, 7921 and 7922.

Query Match 100.0%; Score 194; DB 22; Length 176;
Best Local Similarity 100.0%; Pred. No. 1.8e-19;
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPEELNRYIASLRHYNLVTRQRY 36
|||||
Db 53 YPIKPEAPGEDASPEELNRYIASLRHYNLVTRQRY 88
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RESULT 13

AAH91109
ID AAB91109 standard; Peptide; 36 AA.

AC AAB91109;

XX 22-JUN-2001 (first entry)

DT Parathyroid hormone (PTH) related peptide SEQ ID NO:283.

DE Protection; endogenous therapeutic peptide; peptidase; conjugation;
KW blood component; modification; succinimidyl; maleimido group; amino-
KW hydroxyl; thiol; hormone; growth factor; neurotransmitter.

XX Homo sapiens.

OS Synthetic.

PN WO200069900-A2.

XX 23-NOV-2000.

PF 17-MAY-2000; 2000WO-US13576.

XX 17-MAY-1999; 99US-0134406.

PR 10-SEP-1999; 99US-0153406.

PR 15-OCT-1999; 99US-0159783.

XX (CONJ-) CONJUCHEM INC.

XX Bridon DP, Ezrin AM, Milner PG, Holmes DL, Thibaudeau K;

XX WPI; 2001-112059/12.

XX Modifying and attaching therapeutic peptides to albumin prevents
PT peptidase degradation, useful for increasing length of in vivo activity

PS Disclosure; Page 285; 733pp; English.

XX The present invention describes a modified therapeutic peptide (I)
CC comprising a therapeutically active amino acid region (III) and a
CC reactive group (II) (e.g. succinimidyl and maleimido groups) attached to

CC a less therapeutically active amino acid region (IV), which covalently
CC bonds with amino/hydroxyl/thiol groups on blood components to form a
CC peptidase stabilised therapeutic peptide composed of 3-50 amino acids.
CC (I) are useful for modifying therapeutic peptides e.g. hormones, growth
CC factors and neurotransmitters, to protect them from peptidase activity
CC in vivo for the treatment of various disorders. Endogenous therapeutic
CC peptides are not suitable as drug candidates as they require frequent
CC administration due to rapid degradation by peptidases in the body.
CC Modifying and attaching therapeutic peptides to albumin prevents or
CC reduces the action of peptidases to increase length of activity (half
CC life) and specificity as bonding to large molecules decreases
CC intracellular uptake and interference with physiological processes.
CC AAB90829 to AAB92441 represent peptides which can be used in the
CC exemplification of the present invention.

XX Sequence 36 AA;

Query Match 96.9%; Score 188; DB 22; Length 36;

Best Local Similarity 97.2%; Pred. No. 1.9e-19;

Matches 35; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPEELNRYIASLRHYNLVTRQRY 36
|||||

Db 1 YPIKPEAPGEDASPEELNRYIASLRHYNLVTRQRY 36
|||||

RESULT 14

AAH91226

ID AAB91226 standard; Peptide; 36 AA.

XX AAB91226;

XX 22-JUN-2001 (first entry)

DT Peptide YY SEQ ID NO:400.

DE Protection; endogenous therapeutic peptide; peptidase; conjugation;
KW blood component; modification; succinimidyl; maleimido group; amino-
KW hydroxyl; thiol; hormone; growth factor; neurotransmitter.

XX Homo sapiens.

OS Synthetic.

XX WO200069900-A2.

PN 23-NOV-2000.

XX 17-MAY-2000; 2000WO-US13576.

XX 17-MAY-1999; 99US-0134406.

PR 10-SEP-1999; 99US-0153406.

PR 15-OCT-1999; 99US-0159783.

XX (CONJ-) CONJUCHEM INC.

XX Bridon DP, Ezrin AM, Milner PG, Holmes DL, Thibaudeau K;

XX WPI; 2001-112059/12.

XX Modifying and attaching therapeutic peptides to albumin prevents
PT peptidase degradation, useful for increasing length of in vivo activity

PS Disclosure; Page 329; 733pp; English.

XX The present invention describes a modified therapeutic peptide (I)
CC comprising a therapeutically active amino acid region (III) and a
CC reactive group (II) (e.g. succinimidyl and maleimido groups) attached to
CC a less therapeutically active amino acid region (IV), which covalently
CC bonds with amino/hydroxyl/thiol groups on blood components to form a
CC peptidase stabilised therapeutic peptide composed of 3-50 amino acids.
CC (I) are useful for modifying therapeutic peptides e.g. hormones, growth
CC factors and neurotransmitters, to protect them from peptidase activity

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OM protein - protein search, using sw model

Run on: February 27, 2003, 14:35:44 ; Search time:14 Seconds
(without alignments)
75.659 Million cell updates/sec

Title: US-09-634-363-2
Perfect score: 194
Sequence: 1 YPIKPEAPGEDASPEELNRYASLRHYNLVTRQRY 36

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 262574 seqs, 29422922 residues

Total number of hits satisfying chosen parameters: 262574

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_AA:*

1: /cgn2_6/ptodata/1/1aa/5A_COMB.pep:*
2: /cgn2_6/ptodata/1/1aa/5B_COMB.pep:*
3: /cgn2_6/ptodata/1/1aa/6A_COMB.pep:*
4: /cgn2_6/ptodata/1/1aa/6B_COMB.pep:*
5: /cgn2_6/ptodata/1/1aa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/1/1aa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	%	Query Match	Length	ID	Description
1	194	100.0	36	1	US-08-338-395-2	Sequence 2, Appli
2	194	100.0	36	1	US-08-329-151-2	Sequence 2, Appli
3	194	100.0	36	3	US-09-054-393-2	Sequence 2, Appli
4	194	100.0	36	3	US-09-047-9868-2	Sequence 2, Appli
5	194	100.0	36	4	US-09-229-900-2	Sequence 2, Appli
6	194	100.0	36	5	PCT-US95-14303-2	Sequence 2, Appli
7	184	94.8	36	1	US-07-882-923-3	Sequence 3, Appli
8	184	94.8	36	1	US-08-338-395-1	Sequence 1, Appli
9	184	94.8	36	1	US-08-329-151-1	Sequence 1, Appli
10	184	94.8	36	3	US-09-047-9868-1	Sequence 1, Appli
11	184	94.8	36	4	US-09-181-941-4	Sequence 4, Appli
12	184	94.8	36	5	PCT-US95-14303-1	Sequence 1, Appli
13	175	90.2	36	1	US-08-329-151-9	Sequence 9, Appli
14	162	83.5	36	4	US-09-181-941-1	Sequence 1, Appli
15	158.5	81.7	35	1	US-07-776-272-30	Sequence 30, Appl
16	153	78.9	36	4	US-09-181-941-2	Sequence 2, Appli
17	146	75.3	36	1	US-07-882-923-1	Sequence 1, Appli
18	146	75.3	36	1	US-08-264-030-1	Sequence 1, Appli
19	146	75.3	36	1	US-08-338-395-4	Sequence 4, Appli
20	146	75.3	36	3	US-08-907-403A-2	Sequence 2, Appli
21	146	75.3	36	4	US-09-181-941-5	Sequence 5, Appli
22	146	75.3	36	5	PCT-US95-14303-4	Sequence 4, Appli
23	144	74.2	36	1	US-07-882-923-2	Sequence 2, Appli
24	144	74.2	36	1	US-08-338-395-3	Sequence 3, Appli
25	144	74.2	36	1	US-08-329-151-24	Sequence 24, Appl
26	144	74.2	36	3	US-08-907-403A-1	Sequence 1, Appli
27	144	74.2	36	5	PCT-US95-14303-3	Sequence 3, Appli

28	144	74.2	97	3	US-09-054-393-1	Sequence 1, Appli
29	144	74.2	97	3	US-08-994-946A-6	Sequence 6, Appli
30	144	74.2	97	4	US-09-229-900-1	Sequence 1, Appli
31	144	74.2	97	4	US-09-291-994-6	Sequence 6, Appli
32	141	72.7	36	4	US-09-181-941-3	Sequence 3, Appli
33	129	66.5	24	3	US-09-054-393-7	Sequence 7, Appli
34	129	66.5	24	4	US-09-229-900-7	Sequence 7, Appli
35	127	65.5	32	4	US-09-125-138-10	Sequence 10, Appli
36	115.5	59.5	31	1	US-07-776-272-23	Sequence 23, Appli
37	106	54.6	36	2	US-08-806-203-1	Sequence 1, Appli
38	103	53.1	36	1	US-07-776-272-18	Sequence 18, Appli
39	101	52.1	23	4	US-09-181-941-6	Sequence 6, Appli
40	97	50.0	28	1	US-08-264-030-3	Sequence 3, Appli
41	96	49.5	28	1	US-08-264-030-5	Sequence 5, Appli
42	95	49.0	28	1	US-08-264-030-10	Sequence 10, Appli
43	88	45.4	20	1	US-07-882-923-11	Sequence 11, Appli
44	86.5	44.6	25	1	US-08-264-030-7	Sequence 7, Appli
45	85	43.8	19	1	US-07-882-923-4	Sequence 4, Appli

ALIGNMENTS

RESULT 1
US-08-338-395-2
; Sequence 2, Application US/08338395
; Patent No. 5574010
; GENERAL INFORMATION:
; APPLICANT: McFadden, David W
; TITLE OF INVENTION: TREATMENT OF PANCREATIC TUMORS WITH
; TITLE OF INVENTION: PEPTIDE YY AND ANALOGS THEREOF
; NUMBER OF SEQUENCES: 5
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: POMS, SMITH, LANDE & ROSE
; STREET: 2029 Century Park East 38th Floor
; CITY: Los Angeles
; STATE: CA
; COUNTRY: USA
; ZIP: 90067
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/338,395
; FILING DATE:
; CLASSIFICATION: 514
; ATTORNEY/AGENT INFORMATION:
; NAME: Oldenkamp, David J
; REGISTRATION NUMBER: 29421
; REFERENCE/DOCKET NUMBER: 107012
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 310-788-5046
; TELEFAX: 310-277-1297
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; TYPE: amino acid
; LENGTH: 36 amino acids
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; ORIGINAL SOURCE:
; ORGANISM: HUMAN PEPTIDE YY
US-08-338-395-2

Query Match 100.0%; Score 194; DB 1; Length 36;
Best Local Similarity 100.0%; Pred. No. 8.7e-22;
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
OY 1 YPIKPEAPGEDASPEELNRYASLRHYNLVTRQRY 36
DB 1 YPIKPEAPGEDASPEELNRYASLRHYNLVTRQRY 36

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RESULT 2
US-08-329-151-2
; Sequence 2, Application US/08329151
; Patent No. 5604203
; GENERAL INFORMATION:
; APPLICANT: Balasubramaniam, A.
; TITLE OF INVENTION: ANALOGS OF PEPTIDE YY AND USES
; TITLE OF INVENTION: THEREOF
; NUMBER OF SEQUENCES: 30
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Fish & Richardson
; STREET: 225 Franklin Street
; CITY: Boston
; STATE: Massachusetts
; COUNTRY: U.S.A.
; ZIP: 02110-2804
; COMPUTER READABLE FORM:
; MEDIUM TYPE: 3.5" Diskette, 1.44 MB
; COMPUTER: IBM PS/2 Model 50z or 55sx
; OPERATING SYSTEM: MS-DOS (Version 5.0)
; SOFTWARE: WordPerfect (Version 5.1)
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/329,151
; FILING DATE:
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/038,534
; FILING DATE: 3/29/93
; APPLICATION NUMBER: 08/109,326
; FILING DATE: 08/19/93
; ATTORNEY/AGENT INFORMATION:
; NAME: Paul T. Clark
; REGISTRATION NUMBER: 30,162
; REFERENCE/DOCKET NUMBER: 00537/105001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (617) 542-5070
; TELEFAX: (617) 542-8906
; TELEX: 200154
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 36
; TYPE: amino acid
; STRANDEDNESS: N/A
; TOPOLOGY: linear
US-08-329-151-2

Query Match 100.0%; Score 194; DB 1; Length 36;
Best Local Similarity 100.0%; Pred. No. 8.7e-22;
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 YPIKPEAGEDASPEELNRYASLRHLYNLVTRQY 36
Db 1 YPIKPEAGEDASPEELNRYASLRHLYNLVTRQY 36

RESULT 3
US-09-054-393-2
; Sequence 2, Application US/09054393
; Patent No. 6017879
; GENERAL INFORMATION:
; APPLICANT: Mutter, Manfred
; APPLICANT: Lacroix, Jean S.
; APPLICANT: Grouzmann, Eric
; TITLE OF INVENTION: Template Associated NPV Y2-Receptor
; TITLE OF INVENTION: Agonists
; NUMBER OF SEQUENCES: 8
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Vinson & Elkins LLP
; STREET: 1455 Pennsylvania Avenue, N.W.
; CITY: Washington
; STATE: D.C.
; COUNTRY: U.S.
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; ZIP: 20004-1008
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/054,393
; FILING DATE:
; CLASSIFICATION:
; ATTORNEY/AGENT INFORMATION:
; NAME: Sanzo, Michael A.
; REGISTRATION NUMBER: 36,912
; REFERENCE/DOCKET NUMBER: BMR350/48000
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (202)639-6585
; TELEFAX: (202)639-6604
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 36 amino acids
; TYPE: amino acid
; STRANDEDNESS: not relevant
; TOPOLOGY: not relevant
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
US-09-054-393-2

Query Match 100.0%; Score 194; DB 3; Length 36;
Best Local Similarity 100.0%; Pred. No. 8.7e-22;
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 YPIKPEAGEDASPEELNRYASLRHLYNLVTRQY 36
Db 1 YPIKPEAGEDASPEELNRYASLRHLYNLVTRQY 36

RESULT 4
US-09-047-986B-2
; Sequence 2, Application US/09047986B
; Patent No. 6046167
; GENERAL INFORMATION:
; APPLICANT: Balasubramanian, Ambikaipakan
; TITLE OF INVENTION: PEPTIDE YY ANALOGS
; NUMBER OF SEQUENCES: 20
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Frost & Jacobs, L.L.P.
; STREET: 2500 PNC Center, 201 East Fifth St.
; CITY: Cincinnati
; STATE: OH
; COUNTRY: USA
; ZIP: 45202-4182
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette, 3.5 inch, 1.44 MB storage
; COMPUTER: IBM compatible
; OPERATING SYSTEM: MS-DOS
; SOFTWARE: Word 97
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/047,986B
; FILING DATE: 25 March 1998
; CLASSIFICATION: 514
; ATTORNEY/AGENT INFORMATION:
; NAME: Kristyne A. Bullock
; REGISTRATION NUMBER: 42,371
; REFERENCE/DOCKET NUMBER: 9183030/508
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (513) 651-6731
; TELEFAX: (513) 651-6981
; TELEX: 21-4396 F&J Cin
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 36 amino acids
; TYPE: amino acid
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; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-09-047-986B-2

Query Match      100.0%; Score 194; DB 3; Length 36;
Best Local Similarity 100.0%; Pred. No. 8.7e-22;
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 YPIKPEAPGEDASPEELNRYASLRHYNLVTRQRY 36
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Db 1 YPIKPEAPGEDASPEELNRYASLRHYNLVTRQRY 36

RESULT 5
US-09-229-900-2
; Sequence 2, Application US/09229900
; Patent No. 6288029
; GENERAL INFORMATION:
; APPLICANT: Mutter, Manfred
; APPLICANT: Lacroix, Jean S.
; APPLICANT: Grouzmann, Eric
; TITLE OF INVENTION: Template Associated NPY Y2-Receptor
; TITLE OF INVENTION: Agonists
; NUMBER OF SEQUENCES: 8
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Vinson & Elkins LLP
; STREET: 1455 Pennsylvania Avenue, N.W.
; CITY: Washington
; STATE: D.C.
; COUNTRY: U.S.
; ZIP: 20004-1008
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/229,900
; FILING DATE:
; CLASSIFICATION:
; ATTORNEY/AGENT INFORMATION:
; NAME: Sanzo, Michael A.
; REGISTRATION NUMBER: 36,912
; REFERENCE/DOCKET NUMBER: BMR350/48000
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (202)639-6585
; TELEFAX: (202)639-6604
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 36 amino acids
; TYPE: amino acid
; STRANDEDNESS: not relevant
; TOPOLOGY: not relevant
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
US-09-229-900-2

Query Match      100.0%; Score 194; DB 4; Length 36;
Best Local Similarity 100.0%; Pred. No. 8.7e-22;
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 YPIKPEAPGEDASPEELNRYASLRHYNLVTRQRY 36
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Db 1 YPIKPEAPGEDASPEELNRYASLRHYNLVTRQRY 36

RESULT 6
PCT-US95-14303-2
; Sequence 2, Application PC/TUS9514303
; GENERAL INFORMATION:
; APPLICANT: McFadden, David W
; TITLE OF INVENTION: TREATMENT OF PANCREATIC TUMORS
; TITLE OF INVENTION: WITH PEPTIDE YY AND ANALOGS THEREOF
; NUMBER OF SEQUENCES: 5
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: POMs, SMITH, LANDE & ROSE
; STREET: 2029 Century Park East 38th Floor
; CITY: Los Angeles
; STATE: CA
; COUNTRY: USA
; ZIP: 90067
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: PCT/US95/14303
; FILING DATE: 03 November 1995
; CLASSIFICATION:
; ATTORNEY/AGENT INFORMATION:
; NAME: Oldenkamp, David J
; REGISTRATION NUMBER: 29421
; REFERENCE/DOCKET NUMBER: 107012F
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 310-788-5046
; TELEFAX: 310-277-1297
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 36 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; ORIGINAL SOURCE:
; ORGANISM: HUMAN PEPTIDE YY
PCT-US95-14303-2

Query Match      100.0%; Score 194; DB 5; Length 36;
Best Local Similarity 100.0%; Pred. No. 8.7e-22;
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 YPIKPEAPGEDASPEELNRYASLRHYNLVTRQRY 36
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Db 1 YPIKPEAPGEDASPEELNRYASLRHYNLVTRQRY 36

RESULT 7
US-07-882-923-3
; Sequence 3, Application US/07882923
; Patent No. 5328899
; GENERAL INFORMATION:
; APPLICANT: Boublik, Jaroslav H.
; APPLICANT: Rivier, Jean E.F.
; APPLICANT: Brown, Marvin R.
; APPLICANT: Scott, Neal A.
; TITLE OF INVENTION: NPY PEPTIDE ANALOGS
; NUMBER OF SEQUENCES: 14
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Fitch, Even, Tabin & Flannery
; STREET: 4250 Executive Square, Suite 510
; CITY: La Jolla
; STATE: CA
; COUNTRY: USA
; ZIP: 92037
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/882,923
; FILING DATE: 19920512
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 07/503,198
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; STREET: 1155 Avenue of the Americas
; CITY: New York
; STATE: NY
; COUNTRY: USA
; ZIP: 10036-2811
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: Windows
; SOFTWARE: FastSeq for Windows Version 2.0b
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/181,941
; FILING DATE: 28-Oct-1998
; CLASSIFICATION: <Unknown>
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/574,701
; FILING DATE: 19-DEC-1995
; APPLICATION NUMBER: FR 95 07831
; FILING DATE: 29-JUN-1995
; ATTORNEY/AGENT INFORMATION:
; NAME: Coruzzi, Laura A
; REGISTRATION NUMBER: 30,742
; REFERENCE/DOCKET NUMBER: 3909-0021-999
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 650-493-4935
; TELEFAX: 650-493-5556
; TELEX: 66141 PENNIE
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 36 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: No. 6440690e
; SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-09-181-941-1

Query Match      83.5%; Score 162; DB 4; Length 36;
Best Local Similarity 77.8%; Pred. No. 4.3e-17;
Matches 28; Conservative 5; Mismatches 3; Indels 5

QY   1 YPIKPEAGDASPEELNRYYASLRHLYNLVTRQY 36
    |||:|||||||:|:|:|||||||
Db   1 YPKPESPGEDASPEEMNKYLALRHLYNLVTRQY 36

RESULT 15
US-07-776-272-30
; Sequence 30, Application US/0776272
; Patent No. 5612454
; GENERAL INFORMATION:
; APPLICANT: Kaminuma, Toshihiro
; APPLICANT: Iida, Toshii
; APPLICANT: Tajima, Masahiro
; TITLE OF INVENTION: Process for Purification of Polypeptide
; NUMBER OF SEQUENCES: 31
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Wegner, Cantor, Mueller & Player
; STREET: 1233 20th St. N.W. P.O. Box 18218
; CITY: Washington
; STATE: District of Columbia
; COUNTRY: United States of America
; ZIP: 20036-8218
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/776,272
; FILING DATE: 19911129
; CLASSIFICATION: 530
; ATTORNEY/AGENT INFORMATION:

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; NAME: Player, William E
; REGISTRATION NUMBER: 31,409
; REFERENCE/DOCKET NUMBER: P-450-23167
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 202-887-0400
; TELEFAX: 202-887-0605
; TELEX: 440706
; INFORMATION FOR SEQ ID NO: 30:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 35 amino acids
; TYPE: AMINO ACID
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: YES
; US-07-776-272-30

Query Match      81.7%; Score 158.5; DB 1; Length 35;
Best Local Similarity 88.9%; Pred. No. 1.4e-16;
Matches 32; Conservative 1; Mismatches 2; Indels 1; Gaps 1;

QY 1 YPIKPEAGEDASPEELNRYIASLRHYNLVTQRQY 36
   || ||||| ||||| || ||||| |||||
Db 1 YPAKPEAGEDASPEELSR-XASLRHYNLVTQRQY 35

Search completed: February 27, 2003, 14:38:41
Job time : 15 secs

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GenCore version 5.1.3
Copyright (c) 1993 - 2003 CompuGen Ltd.

OM protein - protein search, using sw model

Run on: February 27, 2003, 14:36:19 ; Search time 12 Seconds
(without alignments)
113.165 Million cell updates/sec

Title: us-09-634-363-2
Perfect score: 194
Sequence: 1 YPIKPEAPGEDASPEELNRYASLRHLYNLVTRQY 36

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 174566 seqs, 37721826 residues
Total number of hits satisfying chosen parameters: 174566

Minimum DB seq length: 0
Maximum DB seq length: 2000000000
Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published_Applications_AA:*

- 1: /cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep.*
- 2: /cgn2_6/ptodata/1/pubpaa/ECT_NEW_PUB.pep.*
- 3: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep.*
- 4: /cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep.*
- 5: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep.*
- 6: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep.*
- 7: /cgn2_6/ptodata/1/pubpaa/ECTUS_PUBCOMB.pep.*
- 8: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep.*
- 9: /cgn2_6/ptodata/1/pubpaa/US09_NEW_PUB.pep.*
- 10: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep.*
- 11: /cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep.*
- 12: /cgn2_6/ptodata/1/pubpaa/US10_PUBCOMB.pep.*
- 13: /cgn2_6/ptodata/1/pubpaa/US60_NEW_PUB.pep.*
- 14: /cgn2_6/ptodata/1/pubpaa/US60_PUBCOMB.pep.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	194	100.0	36	12	US-10-016-969-2
2	184	94.8	36	12	US-10-038-045-4
3	180	92.8	34	12	US-10-016-969-3
4	162	83.5	36	12	US-10-038-045-1
5	153	78.9	36	12	US-10-038-045-2
6	146	75.3	36	12	US-10-038-045-5
7	144	74.2	36	12	US-10-016-969-4
8	144	74.2	97	9	US-10-002-048A-2
9	141	72.7	36	12	US-10-038-045-3
10	132	68.0	34	12	US-10-016-969-5
11	106	54.6	36	12	US-10-016-969-1
12	102	52.6	95	10	US-09-757-712-2
13	101	52.1	23	12	US-10-038-045-6
14	99	51.0	178	9	US-09-965-528-16
15	89	45.9	85	10	US-09-925-300-1040
16	77	39.7	15	12	US-10-016-969-6
17	67	34.5	33	10	US-09-939-825-27
18	55	28.4	1374	9	US-09-900-425A-2
19	54	27.8	446	10	US-09-925-301-1199

20	53	27.3	98	10	US-09-205-658-58	Sequence 58, Appl
21	53	27.3	98	10	US-09-844-353A-58	Sequence 58, Appl
22	53	27.3	492	10	US-09-764-898-206	Sequence 206, Appl
23	53	27.3	509	10	US-09-205-658-46	Sequence 46, Appl
24	53	27.3	509	10	US-09-844-353A-46	Sequence 46, Appl
25	53	27.3	635	10	US-09-844-353A-101	Sequence 101, Appl
26	52.5	27.1	358	10	US-09-864-761-48087	Sequence 48087, A
27	52	26.8	260	9	US-09-840-243B-11	Sequence 11, Appl
28	52	26.8	260	9	US-09-840-243B-12	Sequence 12, Appl
29	50.5	26.0	269	9	US-09-832-129-65	Sequence 65, Appl
30	50.5	26.0	300	9	US-09-832-129-66	Sequence 66, Appl
31	50.5	26.0	365	9	US-09-832-129-67	Sequence 67, Appl
32	50.5	26.0	742	9	US-10-077-111-11	Sequence 11, Appl
33	50	25.8	84	10	US-09-867-550-1630	Sequence 1630, Ap
34	50	25.8	551	9	US-09-870-759-57	Sequence 57, Appl
35	50	25.8	1203	10	US-09-799-875-5	Sequence 5, Appli
36	50	25.8	2923	10	US-09-788-711A-4	Sequence 4, Appli
37	50	25.8	2956	10	US-09-788-711A-2	Sequence 2, Appli
38	49.5	25.5	187	10	US-09-976-451-7	Sequence 7, Appli
39	49	25.3	269	9	US-09-840-243B-13	Sequence 13, Appl
40	49	25.3	315	9	US-09-808-602-65	Sequence 65, Appl
41	49	25.3	844	10	US-09-813-148-4	Sequence 4, Appli
42	49	25.3	844	10	US-09-810-796-14	Sequence 14, Appl
43	49	25.3	871	9	US-10-128-870-20	Sequence 20, Appl
44	48.5	25.0	489	9	US-10-108-605-39	Sequence 39, Appl
45	48	24.7	963	10	US-09-801-368-74	Sequence 74, Appl

ALIGNMENTS

RESULT 1
US-10-016-969-2
; Sequence 2, Application US/10016969
; Patent No. US20020141985A1
; GENERAL INFORMATION:
; APPLICANT: Amnlin Pharmaceuticals, Inc.
; APPLICANT: Pittner, Richard
; APPLICANT: Young, Andrew
; APPLICANT: Paterniti, James
; TITLE OF INVENTION: Peptide YY and Peptide YY Agonists for the Treatment of Metabo
; FILE REFERENCE: 24001-010
; CURRENT APPLICATION NUMBER: US/10/016,969
; CURRENT FILING DATE: 2001-12-14
; PRIOR APPLICATION NUMBER: US 60/256,216
; PRIOR FILING DATE: 2000-12-15
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 2
; LENGTH: 36
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-016-969-2

Query Match 100.0%; Score 194; DB 12;
Best Local Similarity 100.0%; Pred. No. 2.4e-19;
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 YPIKPEAPGEDASPEELNRYASLRHLYNLVTRQY 36
Db 1 YPIKPEAPGEDASPEELNRYASLRHLYNLVTRQY 36

RESULT 2
US-10-038-045-4
; Sequence 4, Application US/10038045
; Patent No. US20020150964A1
; GENERAL INFORMATION:
; APPLICANT: Mor, Amram
; APPLICANT: Vouldoukis, Ioannis
; APPLICANT: Nicolas, Pierre
; TITLE OF INVENTION: PEPTIDES FOR THE ACTIVATION

[illegible]

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; TITLE OF INVENTION: Disorders
; FILE REFERENCE: 24001-010
; CURRENT APPLICATION NUMBER: US/100/016,969
; CURRENT FILING DATE: 2001-12-14
; PRIOR APPLICATION NUMBER: US 60/256,216
; PRIOR FILING DATE: 2000-12-15
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 4
; LENGTH: 36
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-016-969-4

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RESULT 6
US-10-038-045-5
; Sequence 5, Application US/10038045
; Patent No. US20020150964A1
; GENERAL INFORMATION:
; APPLICANT: Mor, Amram
; Vouldoukis, Ioannis
; Nicolas, Pierre
; TITLE OF INVENTION: PEPTIDES FOR THE ACTIVATION
; OF THE IMMUNE SYSTEM IN HUMANS AND ANIMALS
; NUMBER OF SEQUENCES: 16
; CORRESPONDENCE ADDRESS:
US-10-016-969-4
; CURRENT APPLICATION NUMBER: US/10/016,969
; CURRENT FILING DATE: 2001-12-14
; PRIOR APPLICATION NUMBER: US 60/255,216
; PRIOR FILING DATE: 2000-12-15
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 4
; LENGTH: 36
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-016-969-4

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SOFTWARE: FastSeq for Windows Version 2.0.0
CURRENT FILING DATE: 2001-12-14
PRIOR APPLICATION NUMBER: US 60/256,216
PRIORITY FILING DATE: 2000-12-15
NUMBER OF SEQ ID NOS: 6
SOFTWARE: PatentIn version 3.0
SEQ ID NO 1
LENGTH: 36
TYPE: PRT
ORGANISM: Homo sapiens
US-10-016-969-1

Query Match      54.6%; Score 106; DB 12; Length 36;
Best Local Similarity 48.6%; Pred. No. 1.le-07;
Matches 17; Conservative 11; Mismatches 7; Indels 0; Gaps 0;

QY    2 PIKPEAGDASPEELNRYASYLRHYLNLTQRV 36
       ||| | :||| :| | | | | | | | |
DB    2 PLEPYGDNATPEQAQYAADLRRYNMLTRPV 36

RESULT 12
US-09-757-712-2
Sequence 2, Application US/09757712
Patent No. US20010016339A1
GENERAL INFORMATION:
APPLICANT: SOUTHAN, CHRISTOPHER
TITLE OF INVENTION: NOVEL COMPOUNDS
FILE REFERENCE: GP-30015-C1
CURRENT APPLICATION NUMBER: US/09/757,712
CURRENT FILING DATE: 2001-01-10
PRIOR APPLICATION NUMBER: EP 97307187.1
PRIOR FILING DATE: 1997-09-16
PRIOR APPLICATION NUMBER: 09/110,715
PRIOR FILING DATE: 1998-07-07
NUMBER OF SEQ ID NOS: 2
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 2
LENGTH: 95
TYPE: PRT
ORGANISM: HOMO SAPIENS
US-09-757-712-2

Query Match      52.6%; Score 102; DB 10; Length 95;
Best Local Similarity 45.7%; Pred. No. 1.le-06;
Matches 16; Conservative 11; Mismatches 8; Indels 0; Gaps 0;

QY    2 PIKPEAGDASPEELNRYASYLRHYLNLTQRV 36
       ||| | :||| :| | | | | | | | |
DB    31 PLEPLYGDNTTPEQAQYTAEALRYYINMLTHRV 65

RESULT 13
US-10-038-045-6
Sequence 6, Application US/10038045
Patent No. US20020150964A1
GENERAL INFORMATION:
APPLICANT: Mor, Anram
              Voulcoukis, Ioannis
              Nicolas, Pierre
TITLE OF INVENTION: PEPTIDES FOR THE ACTIVATION
                  OF THE IMMUNE SYSTEM IN HUMANS AND ANIMALS
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: NY
COUNTRY: USA
ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
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; NAME/KEY: misc.feature
; OTHER INFORMATION: Incyte ID No. US20020187523A1 5037143CD1
US-09-965-528-16

Query Match 51.0%; Score 99; DB 9; Length 178;
Best Local Similarity 47.1%; Pred. No. 5.8e-06;
Matches 16; Conservative 11; Mismatches 7; Indels 0; Gaps 0;

QY 2 PIKPEAPGEDASPEELNRYRYASLRHYLNLVTQR 35
Db 31 PLEPVYGDNATPEQMAQYAADLRRYINMLTRPR 64

RESULT 15

US-09-925-300-1040
; Sequence 1040, Application US/09925300
; Patent No. US20020151681A1
; GENERAL INFORMATION:
; APPLICANT: Craig Rosen,
; APPLICANT: Steve Ruben,
; TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
; FILE REFERENCE: PA101
; CURRENT APPLICATION NUMBER: US/09/925,300
; PRIOR FILING DATE: 2001-08-10
; PRIOR APPLICATION NUMBER: PCT/US00/05988
; PRIOR FILING DATE: 2000-03-08
; PRIOR APPLICATION NUMBER: 60/124,270
; PRIOR FILING DATE: 1999-03-12
; NUMBER OF SEQ ID NOS: 1890
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 1040
; LENGTH: 85
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (30)
; OTHER INFORMATION: xaa equals any of the naturally occurring L-amino acids
US-09-925-300-1040

Query Match 45.9%; Score 89; DB 10; Length 85;
Best Local Similarity 66.7%; Pred. No. 5.2e-05;
Matches 14; Conservative 6; Mismatches 1; Indels 0; Gaps 0;

QY 16 ELNRYRYASLRHYLNLVTQR 36
Db 32 DMARYYSALRHYINLITQR 52

Search completed: February 27, 2003, 14:39:00
Job time : 13 secs

Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPELNRYRYASLRHYNLVTRQRY 36
|||||
Db 29 YPIKPEAPGEDASPELNRYRYASLRHYNLVTRQRY 64
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RESULT 3
S34568
peptide YY precursor (clone 11) - human (fragment)
C:Species: Homo sapiens (man)
C>Date: 02-Dec-1993 #sequence_revision 13-Mar-1997 #text_change 09-May-1997
C:Accession: S34568
R:Kohri, K.; Nata, K.; Yonekura, H.; Nagai, A.; Konno, K.; Okamoto, H.
Biochim. Biophys. Acta 1173, 345-349, 1993
A:Title: Cloning and structural determination of human peptide YY cDNA and gene.
A:Reference number: S33795; MUID:93305732; PMID:8318545
A:Accession: S34568
A:Status: preliminary
A:Molecule type: mRNA
A:Residues: 1-90 <KOH>
C:Superfamily: pancreatic hormone

Query Match 100.0%; Score 194; DB 2; Length 90;
Best Local Similarity 100.0%; Pred. No. 1.4e-18;
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPELNRYRYASLRHYNLVTRQRY 36
|||||
Db 29 YPIKPEAPGEDASPELNRYRYASLRHYNLVTRQRY 64
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RESULT 4
S33795
peptide YY (clone S) - human
C:Species: Homo sapiens (man)
C>Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 20-Jun-2000
C:Accession: S33795
R:Kohri, K.; Nata, K.; Yonekura, H.; Nagai, A.; Konno, K.; Okamoto, H.
Biochim. Biophys. Acta 1173, 345-349, 1993
A:Title: Cloning and structural determination of human peptide YY cDNA and gene.
A:Reference number: S33795; MUID:93305732; PMID:8318545
A:Accession: S33795
A:Status: preliminary
A:Molecule type: mRNA
A:Residues: 1-97 <KOH>
A:Cross-references: GB:DJ3897; NID:g391723; PIDN:BAA02997.1; PTD:g391724
C:Superfamily: pancreatic hormone

Query Match 100.0%; Score 194; DB 2; Length 97;
Best Local Similarity 100.0%; Pred. No. 1.5e-18;
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPELNRYRYASLRHYNLVTRQRY 36
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Db 29 YPIKPEAPGEDASPELNRYRYASLRHYNLVTRQRY 64
|||||

RESULT 5
YYPG
peptide YY - pig
C:Species: Sus scrofa domestica (domestic pig)
C>Date: 30-Apr-1982 #sequence_revision 30-Apr-1982 #text_change 23-Aug-1996
C:Accession: A01574
R:Fatemoto, K.
Proc. Natl. Acad. Sci. U.S.A. 79, 2514-2518, 1982
A:Title: Isolation and characterization of peptide YY (pYY), a candidate gut hormone the
A:Reference number: A01574; MUID:82222168; PMID:6953409
A:Accession: A01574
A:Molecule type: protein
A:Residues: 1-36 <FAT>
C:Superfamily: pancreatic hormone
C:Keywords: amidated carboxyl end; hormone

F.36/Modified site: amidated carboxyl end (Tyr) #status experimental

Query Match 94.8%; Score 184; DB 1; Length 36;
Best Local Similarity 94.4%; Pred. No. 1e-17;
Matches 34; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPELNRYRYASLRHYNLVTRQRY 36
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Db 1 YPAKPEAPGEDASPELSRYASLRHYNLVTRQRY 36
|||||

RESULT 6
A60416
peptide YY - dog
C:Species: Canis lupus familiaris (dog)
C>Date: 11-Feb-1993 #sequence_revision 11-Feb-1993 #text_change 17-Mar-1999
C:Accession: A60416
R:Eysselein, V.E.; Eberlein, G.A.; Grandt, D.; Schaeffer, M.; Zehres, B.; Behn, U.; S
Peptides 11, 111-116, 1990
A:Title: Structural characterization of canine PYY.
A:Reference number: A60416; MUID:90259843; PMID:2342986
A:Accession: A60416
A:Molecule type: protein
A:Residues: 1-36 <EYS>
C:Superfamily: pancreatic hormone
C:Keywords: amidated carboxyl end; hormone; intestine
F.36/Modified site: amidated carboxyl end (Tyr) #status experimental

Query Match 94.8%; Score 184; DB 2; Length 36;
Best Local Similarity 94.4%; Pred. No. 1e-17;
Matches 34; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPELNRYRYASLRHYNLVTRQRY 36
|||||
Db 1 YPAKPEAPGEDASPELSRYASLRHYNLVTRQRY 36
|||||

RESULT 7
A29364
peptide YY precursor - rat
C:Species: Rattus norvegicus (Norway rat)
C>Date: 31-Dec-1988 #sequence_revision 31-Dec-1988 #text_change 16-Jul-1999
C:Accession: A37955; A29364; JT0416
R:Krasinski, S.D.; Wheeler, M.B.; Leiter, A.B.
Mol. Endocrinol. 5, 433-440, 1991
A:Title: Isolation, characterization, and developmental expression of the rat peptide
A:Reference number: A37955; MUID:91367188; PMID:1890992
A:Accession: A37955
A:Status: preliminary
A:Molecule type: DNA
A:Residues: 1-98 <KRA>
A:Cross-references: GB:S57220; NID:g235283; PIDN:AA819752.1; PID:g235284
R:Leiter, A.B.; Toder, A.; Wolfe, H.J.; Taylor, I.L.; Cooperman, S.; Mandel, G.; Good
J. Biol. Chem. 262, 12984-12988, 1987
A:Title: Peptide YY. Structure of the precursor and expression in exocrine pancreas.
A:Reference number: A29364; MUID:88007492; PMID:3654598
A:Accession: A29364
A:Molecule type: mRNA
A:Residues: 1-98 <LEI>
A:Cross-references: GB:M17523; NID:g204316; PIDN:AAA41222.1; PID:g204317
R:Corder, R.; Gaillard, R.C.; Boehlen, P.
Regul. Pept. 21, 253-261, 1988
A:Title: Isolation and sequence of rat peptide YY and neuro peptide Y.
A:Reference number: JT0416; MUID:88321122; PMID:3413293
A:Accession: JT0416
A:Molecule type: protein
A:Residues: 29-64 <COR>
A:Experimental source: colon
C:Superfamily: pancreatic hormone

Query Match 94.8%; Score 184; DB 2; Length 98;
Best Local Similarity 94.4%; Pred. No. 3.2e-17;
Matches 34; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Query Match 79.9%; Score 155; DB 2; Length 36;
Best Local Similarity 75.0%; Pred. No. 7e-14;
Matches 27; Conservative 4; Mismatches 5; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPEELNRYASLRHYLNLVTRQY 36
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Db 1 YPKPENPGEDAPPEELAKYVYALRHYINLITRQY 36

RESULT 13
S27054
neuropeptide Y - Atlantic cod
C:Species: Gadus morhua (Atlantic cod)
C>Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 31-Oct-1997
C:Accession: S27054
R:Jensen, J.; Conlon, J.M.
Eur. J. Biochem. 210, 405-410, 1992
A:Title: Characterization of peptides related to neuropeptide tyrosine and peptide tyros
A:Reference number: S27054; MUID:93092973; PMID:1459125
A:Accession: S27054
A:Status: preliminary
A:Molecule type: protein
A:Residues: 1-36 <JEN>
C:Superfamily: pancreatic hormone

Query Match 78.4%; Score 152; DB 2; Length 36;
Best Local Similarity 72.2%; Pred. No. 1.8e-13;
Matches 26; Conservative 6; Mismatches 4; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPEELNRYASLRHYLNLVTRQY 36
||||| ||||| :|| :||||:|||||
Db 1 YPIKPEAPGEDAPPEELAKYVYALRHYINLITRQY 36

RESULT 14
S26954
peptide YY-related protein, intestinal - chicken
N:Alternate names: neuropeptide Y homolog; peptide tyrosine-tyrosine-related protein
C:Species: Gallus gallus (Chicken)
C>Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 07-Feb-1997
C:Accession: S26954
R:Conlon, J.M.; O'Harte, F.
FEBS Lett. 313, 225-228, 1992
A:Title: The primary structure of a PYX-related peptide from chicken intestine suggests
A:Reference number: S26954; MUID:93076900; PMID:1446739
A:Accession: S26954
A:Molecule type: protein
A:Residues: 1-37 <CON>
C:Superfamily: pancreatic hormone
C:Keywords: amidated carboxyl end; hormone; intestine; neuropeptide

Query Match 75.8%; Score 147; DB 2; Length 37;
Best Local Similarity 69.4%; Pred. No. 8.3e-13;
Matches 25; Conservative 8; Mismatches 3; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPEELNRYASLRHYLNLVTRQY 36
|| ||| ||||| :||||:|||||
Db 2 YPKPENPGEDAPPEELAKYVYALRHYINLITRQY 37

RESULT 15
A41979
neuropeptide Y precursor - chicken
C:Species: Gallus gallus (chicken)
C>Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 21-Jul-2000
C:Accession: A41979
R:Blomqvist, A.G.; Soderberg, C.; Lundell, I.; Milner, R.J.; Larhammar, D.
Proc. Natl. Acad. Sci. U.S.A. 89, 2350-2354, 1992
A:Title: Strong evolutionary conservation of neuropeptide Y: sequences of chicken, goldf
A:Reference number: A41979; MUID:92196116; PMID:1549597
A:Accession: A41979
A:Status: preliminary
A:Molecule type: mRNA

A:Residues: 1-97 <BLO>
A:Cross-references: GB:M87294; NID:g212458; PIDN:AAA48991.1; PID:g212459
A:Experimental source: central nervous system
A>Note: sequence extracted from NCBI backbone (NCBIP:88404)
C:Function:
A:Description: neuropeptide inducing a number of behavioral effects including stimuli
C:Superfamily: pancreatic hormone
C:Keywords: amidated carboxyl end; appetite; hormone; neuropeptide
F:1-28/Domain: signal sequence #status predicted <Sig>
F:29-64/Product: neuropeptide Y #status predicted <MAT>
F:65-97/Domain: carboxyl-terminal propeptide #status predicted <CTP>
F:64/Modified site: amidated carboxyl end (Tyr) (amide in mature form from following

Query Match 75.8%; Score 147; DB 2; Length 97;
Best Local Similarity 66.7%; Pred. No. 2.5e-12;
Matches 24; Conservative 8; Mismatches 4; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPEELNRYASLRHYLNLVTRQY 36
|| ||| ||||| :|| :||||:|||||
Db 29 YPKPENPGEDAPPEELAKYVYALRHYINLITRQY 64

Search completed: February 27, 2003, 14:38:20
Job time : 46 secs

GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.

OM protein - protein search, using sw model

Run on: February 27, 2003, 14:16:23 ; Search time 29 Seconds
(without alignments)

51.488 Million cell updates/sec

Title: US-09-634-363-2

Perfect score: 194

Sequence: 1 YPIKPEAPGEDASPELNRYASLRHYNLVTRORY 36

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 112892 seqs, 41476328 residues

Total number of hits satisfying chosen parameters: 112892

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : SwissProt_40.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	194	100.0	97	1 PYV_HUMAN	P10082 homo sapien
2	184	94.8	36	1 PYV_PIG	P01305 sus scrofa
3	184	94.8	93	1 PYV_MOUSE	Q9eps2 mus musculu
4	184	94.8	98	1 PYV_RAT	P10631 rattus norv
5	168	86.6	97	1 PYV_BOVIN	P51694 bos taurus
6	162	83.5	36	1 SPV_PHYBI	P80952 phyllomedus
7	158	81.4	36	1 PYV_AMICA	P29205 amia calva
8	156	80.4	36	1 PYV_LEPSP	P09473 lepisosteus
9	155	79.9	36	1 PYV_ONCKI	P09474 oncorhynch
10	155	79.9	36	1 PYV_RAJRH	P29206 raja rhina
11	154	79.4	97	1 PYV_BRARE	Q918p2 brachydanio
12	154	79.4	99	1 NEUY_DICLA	Q9pta0 dicentrarch
13	153	78.9	36	1 NEUY_ONCMY	P29071 oncorhynch
14	153	78.9	36	1 PYV_RANRI	P29204 rana ridibu
15	152	78.4	36	1 NEUY_GADMO	P80167 gadus morhu
16	149	76.8	95	1 NEUY_ICTPU	Q919d3 ictalurus p
17	147	75.8	37	1 PYV_CHICK	P29203 gallus gall
18	147	75.8	96	1 NEUY_BRARE	Q918p3 brachydanio
19	147	75.8	97	1 NEUY_CHICK	P28673 gallus gall
20	146	75.3	36	1 NEUY_PIG	P01304 sus scrofa
21	144	74.2	36	1 NEUY_RABIT	P09640 oryctolagus
22	144	74.2	36	1 PYV_ORENI	P81028 oreochromis
23	144	74.2	97	1 NEUY_HUMAN	P01303 homo sapien
24	144	74.2	97	1 NEUY_MOUSE	P57774 mus musculu
25	144	74.2	98	1 NEUY_RAT	P07808 rattus norv
26	143	73.7	36	1 NEUY_SHEEP	P14765 ovis aries
27	141	72.7	36	1 NEUY_RANRI	P29949 rana ridibu
28	141	72.7	97	1 NEUY_XENLA	P33689 xenopus lae
29	140	72.2	96	1 NEUY_CARAU	P28672 carassius a
30	139	71.6	97	1 PY_DICLA	Q9pt98 dicentrarch
31	137	70.6	99	1 PYV_DICLA	Q9pt99 dicentrarch
32	136	70.1	98	1 NEUY_TORMA	P28674 torpedo mar
33	135	69.6	93	1 PYV_LAMFL	P48098 lampetra fl

RESULT 1

ID	PYV_HUMAN	STANDARD;	PRT;	97 AA.
AC	P10082;			
DT	01-MAR-1989 (Rel. 10, Created)			
DT	01-NOV-1995 (Rel. 32, Last sequence update)			
DT	16-OCT-2001 (Rel. 40, Last annotation update)			
DE	Peptide YY precursor (PYV) (Peptide tyrosine tyrosine).			
GN	PYY.			
OS	Homo sapiens (Human).			
OC	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;			
OC	Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.			
OX	NCBI_TaxID:9606;			
RN	[1]			
RP	SEQUENCE FROM N.A.			
RC	TISSUE=Colon mucosa;			
RX	MEDLINE=93305732; PubMed=8318545;			
RA	Kohri K., Nata K., Yonekura H., Nagai A., Konno K., Okamoto H.;			
RT	"Cloning and structural determination of human peptide YY cDNA and			
RL	gene.";			
RL	Biochim. Biophys. Acta 1173:345-349(1993).			
RN	[2]			
RP	SEQUENCE FROM N.A.			
RC	TISSUE=Lymphocytes;			
Herzog H.;				
RN	Submitted (NOV-1993) to the EMBL/GenBank/DBJ databases.			
[3]				
RX	SEQUENCE OF 29-64, AND SYNTHESIS OF 29-64.			
RP	MEDLINE=89076307; PubMed=3202875;			
RA	Tatemoto K., Nakano I., Makk G., Angwin P., Mann M., Schilling J.,			
Go V.L.W.;				
RT	"Isolation and primary structure of human peptide YY.";			
RL	Biochem. Biophys. Res. Commun. 157:713-717(1988).			
RN	[4]			
RP	SEQUENCE OF 29-64.			
RX	MEDLINE=90068171; PubMed=2587421;			
RA	Eberlein G.A., Eysselein V.E., Schaeffer M., Layer P., Grandt D.,			
Goebell H., Niebel W., Davis M., Lee T.D., Shively J.E.,				
Reeve J.R. Jr.;				
RT	"A new molecular form of PYY: structural characterization of human			
PYY(3-36) and PYY(1-36).";				
RL	Peptides 10:797-803(1989).			
CC	-!- FUNCTION: THIS GUT PEPTIDE INHIBITS EXOCRINE PANCREATIC SECRETION,			
	HAS A VASOCONSTRICTORY ACTION AND INHIBITS JEJUNAL AND COLONIC			
	MOBILITY.			
CC	-!- SUBCELLULAR LOCATION: Secreted.			
CC	-!- ALTERNATIVE PRODUCTS: 2 ISOFORMS: A LONG FORM (SHOWN HERE) AND A			
CC	SHORT FORM; ARE PRODUCED BY ALTERNATIVE SPLICING.			
CC	-!- SIMILARITY: BELONGS TO THE NPY / PYY / PYY FAMILY.			
CC	-----			
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CC	between the Swiss Institute of Bioinformatics and the EMBL outstation -			
CC	the European Bioinformatics Institute. There are no restrictions on its			
CC	use by non-profit institutions as long as its content is in no way			
CC	modified and this statement is not removed. Usage by and for commercial			
CC	entities requires a license agreement (See http://www.isb-sib.ch/announce/)			

ALIGNMENTS

34	133	68.6	36	1	PYV_MYOSC	P09641 myoxocephal
35	133	68.6	104	1	NEUY_LAMFL	P48097 lampetra fl
36	132	68.0	69	1	PYV_LOPAM	P09475 lophius ame
37	122	62.9	36	1	PYV_PETMA	P80024 petromyzon
38	113	58.2	131	1	PAHO_BOVIN	P01302 bos taurus
39	111	57.2	36	1	PAHO_PIG	P01300 sus scrofa
40	111	57.2	93	1	PAHO_CANFA	P01299 canis famil
41	109	56.2	36	1	PAHO_CERSI	P37999 ceratotheri
42	107	55.2	36	1	PAHO_LARAR	P41337 larus argen
43	106	54.6	36	1	PAHO_MACMU	P33684 macaca mula
44	106	54.6	36	1	PAHO_RABIT	P41336 oryctolagus
45	106	54.6	36	1	PAHO_TAPPI	P39659 tapirus pin

```

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CC EMBL; D13897; BAA02997.1; -
CC EMBL; D13897; BAA02998.1; -
DR EMBL; D13897; BAA02998.1; -
DR EMBL; D13899; BAA03000.1; -
DR EMBL; D13902; BAA03002.1; -
DR EMBL; L25648; AAA36433.1; -
DR PIR; A31358; A31358.
DR PIR; A60676; A60676.
DR HSSP; P01303; IRON.
DR HSSP; HGNC:9748; PYY.
DR MIM; 600781; -.
DR InterPro; IPR001955; Pancreatic_hormn.
DR Pfam; PF00159; hormone3; 1.
DR PRINTS; PR00278; PANCHORMONE.
DR ProDom; PD001267; Pancreatic_hormn; 1.
DR SMART; SM00309; PAH; 1.
DR PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.
DR PROSITE; PS0276; PANCREATIC_HORMONE_2; 1.
KW Hormone; Cleavage on pair of basic residues; Pancreas; Amidation;
KW Signal; Alternative splicing.
FT SIGNAL 1 28
FT PEPTIDE 29 64 PEPTIDE YY.
FT PROPEP 68 97
FT MOD_RES 64 64
FT VARSPIC 91 97
FT VARIANT 72 72
FT FTID-VAR_006382.
FT SEQUENCE 97 AA; 11046 MW; DD16B73407F656A4 CRC64;
Query Match 100.0%; Score 194; DB 1; Length 97;
Best Local Similarity 100.0%; Pred. No. 1.1e-19;
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPELNRYASLRHLYNLVTRQRY 36
Db 29 YPIKPEAPGEDASPELNRYASLRHLYNLVTRQRY 64

RESULT 2
ID PYY_PIG STANDARD; PRT; 36 AA.
AC P01305;
DC 21-JUL-1986 (Rel. 01, Created)
DT 21-JUL-1986 (Rel. 01, Last sequence update)
DE 16-OCT-2001 (Rel. 40, Last annotation update)
DE Peptide YY (PYY) (Peptide tyrosine tyrosine).
GN PYY.
OS Sus scrofa (Pig), and
OS Canis familiaris (Dog).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae; Sus.
OX NCBI_TaxID=9823, 9615;
RN [1]
RP SPECIES=Pig;
RC MEDLINE=8222168; PubMed=6953409;
RX Tatemoto K.;
RA "Isolation and characterization of peptide YY (PYY), a candidate gut
RT hormone that inhibits pancreatic exocrine secretion.";
RL Proc. Natl. Acad. Sci. U.S.A. 79:2514-2518(1982).
RN [2]
RP SPECIES=C.familiaris;
RC MEDLINE=90259843; PubMed=2342986;
RX Eysselein V.E., Eberlein G.A., Grandt D., Schaeffer M., Zehres B.,
RA Meyer U., Schaefer D., Goebell H., Davis M., Lee T.D., Shively J.E.,
RA Meyer H.E., Reeve J.R. Jr.;
RT "Structural characterization of canine PYY.";
RL Peptides 11:111-116(1990).
CC -!- FUNCTION: THIS GUT PEPTIDE INHIBITS EXOCRINE PANCREATIC SECRETION,
CC HAS A VASOCONSTRICTORY ACTION AND INHIBITS JEJUNAL AND COLONIC
CC MOBILITY.

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CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- SIMILARITY: BELONGS TO THE NPY / PYY / PYY FAMILY.
DR PIR; A01574; YYPG.
DR PIR; A60416; A60416.
DR HSSP; P01303; IRON.
DR InterPro; IPR001955; Pancreatic_hormn.
DR Pfam; PF00159; hormone3; 1.
DR PRINTS; PR00278; PANCHORMONE.
DR ProDom; PD001267; Pancreatic_hormn; 1.
DR SMART; SM00309; PAH; 1.
DR PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.
DR PROSITE; PS0276; PANCREATIC_HORMONE_2; 1.
KW Hormone; Amidation.
FT MOD_RES 36 36 AMIDATION.
FT SEQUENCE 36 AA; 4242 MW; 02CD6B8C586DCC8D CRC64;
Query Match 94.8%; Score 184; DB 1; Length 36;
Best Local Similarity 94.4%; Pred. No. 7.7e-19;
Matches 34; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPELNRYASLRHLYNLVTRQRY 36
Db 1 YPAKPEAPGEDASPELSRYASLRHLYNLVTRQRY 36

RESULT 3
ID PYY_MOUSE STANDARD; PRT; 93 AA.
AC Q9EPS2;
DC 16-OCT-2001 (Rel. 40, Created)
DT 16-OCT-2001 (Rel. 40, Last sequence update)
DT 15-JUN-2002 (Rel. 41, Last annotation update)
DE Peptide YY precursor (PYY) (Peptide tyrosine tyrosine) (Fragment).
GN PYY.
OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX NCBI_TaxID=10090;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=BALB/c; TISSUE=Liver;
RA Brown G.J., James R., Eddie L.W.;
RL Submitted (DEC-2000) to the EMBL/GenBank/DBJ databases.
CC -!- FUNCTION: THIS GUT PEPTIDE INHIBITS EXOCRINE PANCREATIC SECRETION,
CC HAS A VASOCONSTRICTORY ACTION AND INHIBITS JEJUNAL AND COLONIC
CC MOBILITY.
CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- SIMILARITY: BELONGS TO THE NPY / PYY / PYY FAMILY.
-----
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-----
CC EMBL; AF325866; AAG42908.1; -.
DR HSSP; P01303; IRON.
DR MGD; MGI:99924; PYY.
DR InterPro; IPR001955; Pancreatic_hormn.
DR Pfam; PF00159; hormone3; 1.
DR PRINTS; PR00278; PANCHORMONE.
DR ProDom; PD001267; Pancreatic_hormn; 1.
DR SMART; SM00309; PAH; 1.
DR PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.
DR PROSITE; PS0276; PANCREATIC_HORMONE_2; 1.
KW Hormone; Cleavage on pair of basic residues; Pancreas; Amidation;
KW Signal.
FT SIGNAL 1 28 BY SIMILARITY.
FT PEPTIDE 29 64 PEPTIDE YY.
FT PROPEP 68 93
FT MOD_RES 64 64
FT AMIDATION (G-65 PROVIDE AMIDE GROUP).
FT MISSING (IN SHORT ISOFORM).
FT T -> R.
FT FTID-VAR_006382.
FT SEQUENCE 97 AA; 11046 MW; DD16B73407F656A4 CRC64;
Query Match 100.0%; Score 194; DB 1; Length 97;
Best Local Similarity 100.0%; Pred. No. 1.1e-19;
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPELNRYASLRHLYNLVTRQRY 36
Db 29 YPIKPEAPGEDASPELNRYASLRHLYNLVTRQRY 64

RESULT 2
ID PYY_PIG STANDARD; PRT; 36 AA.
AC P01305;
DC 21-JUL-1986 (Rel. 01, Created)
DT 21-JUL-1986 (Rel. 01, Last sequence update)
DE 16-OCT-2001 (Rel. 40, Last annotation update)
DE Peptide YY (PYY) (Peptide tyrosine tyrosine).
GN PYY.
OS Sus scrofa (Pig), and
OS Canis familiaris (Dog).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae; Sus.
OX NCBI_TaxID=9823, 9615;
RN [1]
RP SPECIES=Pig;
RC MEDLINE=8222168; PubMed=6953409;
RX Tatemoto K.;
RA "Isolation and characterization of peptide YY (PYY), a candidate gut
RT hormone that inhibits pancreatic exocrine secretion.";
RL Proc. Natl. Acad. Sci. U.S.A. 79:2514-2518(1982).
RN [2]
RP SPECIES=C.familiaris;
RC MEDLINE=90259843; PubMed=2342986;
RX Eysselein V.E., Eberlein G.A., Grandt D., Schaeffer M., Zehres B.,
RA Meyer U., Schaefer D., Goebell H., Davis M., Lee T.D., Shively J.E.,
RA Meyer H.E., Reeve J.R. Jr.;
RT "Structural characterization of canine PYY.";
RL Peptides 11:111-116(1990).
CC -!- FUNCTION: THIS GUT PEPTIDE INHIBITS EXOCRINE PANCREATIC SECRETION,
CC HAS A VASOCONSTRICTORY ACTION AND INHIBITS JEJUNAL AND COLONIC
CC MOBILITY.

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KW Signal.
FT SIGNAL      1    28
FT PEPTIDE     29    64
FT PROPEP     60    98
FT MOD_RES     64    68
SQ SEQUENCE    98 AA;  1121 MW;   994C0C3A06A8A7DE CRC64;

Query Match          94.8%; Score 184; DB 1; Length 98;
Best Local Similarity 94.4%; Pred. No. 2.6e-18;
Matches 34; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPEELNRYYSRLHYLNLVTRORY 36
   ||| | | | | | | | | | : | | | | | | | | | | | | | | | |
DB 29 YPAKPEAPGEDASPEELSRYYSRLHYLNLVTRORY 64

RESULT 5
PPY_BOVIN STANDARD; PRT; 97 AA.
AC P51694;
DT 01-OCT-1996 (Rel. 34, Created)
DD 01-OCT-1996 (Rel. 34, Last sequence update)
DE 16-OCT-2001 (Rel. 40, Last annotation update)
DEF Peptide YY precursor (PPY) (peptide tyrosine tyrosine).
GN PPY.
OS Bos taurus (Bovine).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae;
OC Bovidae; Bovinae; Bos.
OX NCBI_TaxID=9913;
RN [1]
RS SEQUENCE FROM N.A.
RX MEDLINE=95132646; PubMed=7831336;
RA Herzog H., Hort Y., Schneider R., Shine J.;
RT "Seminalplasmin: recent evolution of another member of the
   neuropeptide Y gene family.";
RL Proc. Natl. Acad. Sci. U.S.A. 92:594-598(1995).
CC -1- FUNCTION: THIS GUT PEPTIDE INHIBITS EXOCRINE PANCREATIC SECRETION,
   HAS A VASOCONSTRICTORY ACTION AND INHIBITS JEJUNAL AND COLONIC
   MOBILITY.
CC -1- SUBCELLULAR LOCATION: Secreted.
CC -1- SIMILARITY: BELONGS TO THE NPY / PPY / PYV FAMILY.
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or send an email to licensee@isb-sib.ch).
-----
EMBL; L37369; AAC37326.1; ..
DR HSPP; P01303; IRON.
DR InterPro; IPR001955; Pancreatic_hormn.
DR Pfam; PF00159; hormone3; 1.
DR PRINTS; PR00278; PANCHORMONE.
DR ProDom; PD001267; Pancretatic_hormn; 1.
DR SMART; SM00309; PAH; 1.
DR PROSITE; PS00285; PANCREATIC_HORMONE_1; 1.
DR PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.
DK Hormone; Cleavage on pair of basic residues; Pancreas; Amidation;
Signal.
FT SIGNAL      1    28
FT PEPTIDE     29    64
FT PROPEP     68    97
FT MOD_RES     64    64
SQ SEQUENCE    97 AA;  11092 MW;   B3A7BGAT6GBBB3AE0 CRC64;

Query Match          86.6%; Score 168; DB 1; Length 97;
Best Local Similarity 83.3%; Pred. No. 3.8e-16;
Matches 30; Conservative 3; Mismatches 3; Indels 0; Gaps 0;

OV 1 YPIKPEAPGEDASPEELNRYYSRLHYLNLVTRORY 36

```

	TISSUE=Pancreas;
RC	MEDLINE=91296574; PubMed=2067973;
RX	Conlon J.M., Bjénning C., Moon T.W., Youson J.H., Thim L.;
RA	"Neuropeptide Y-related peptides from the pancreas of a teleostean
RT	(eel), holosteian (bowfin) and elasmobranch (skate) fish.";
RL	Peptides 12:221-226(1991).
CC	-I- SUBCELLULAR LOCATION: Secreted.
DD	SIMILARITY: BELONGS TO THE NPY / PPY / PYF FAMILY.
HSPG	P01303; IRON.
DR	InterPro: IPRO01955; Pancreat_hormn.
DR	Pfam: PF00159; hormone3_1.
DR	PRINTS: PR00278; PANCORMONE.
DR	PRODOM: PD001267; Pancreat_hormn; 1.
DR	SMART: SMO0309; PAH; 1.
DR	PROSITE: PS00265; PANCREATIC_HORMONE_1; 1.
DR	PROSITE: PS00276; PANCREATIC_HORMONE_2; 1.
KW	Hormone; Amidation.
FT	MOD_RES 36 36 AMIDATION.
SQ	SEQUENCE 36 AA; 4333 MW; 56B46F3C08666671 CRC64;
	Query Match 81.4%; Score 158; DB 1; Length 36;
	Best Local Similarity 77.8%; Pred No. 2.7e-15;
	Matches 28; Conservative 3; Mismatches 5; Indels 0; Gaps
OY	1 YPKPEAPGEDASPEELNRYASLRHYLNLVTRQR 36 : :
Dd	1 YPPKPNPGEDAPEELARYTALRHYINLTIRQR 36 :
RESULT 8	
PYY_LEPSP	STANDARD; PRT; 36 AA.
ID	PYY_LEPSP
PC	P09473;
DT	01-MAR-1989 (Rel. 10, Created)
DT	01-MAR-1989 (Rel. 10, Last sequence update)
DE	Peptide YY-like (PYY) (Neuropeptide Y-related peptide).
OS	Lepisosteus spatula (Alligator gar) (Atractosteus spatula),
OS	Scyliorhinus canicula (Spotted dogfish) (Spottet catshark), and
OS	Squalus acanthias (Spiny dogfish).
OC	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC	Actinopterygii; Neopterygii; Semionotiformes; Lepisosteidae;
OX	Lepisosteus.
NCBI_TaxId	7917, 7830, 7797;
[1]	
SEQUENCE.	
SPECIES=L.spatula; TISSUE=Pancreeas;	
MEDLINE=88030594; Pubmed=3311873;	
Pollock H.G., Kimmel J.R., Hamilton J.W., Rouse J.B., Ebner K.E.,	
Lance V., Rawitch A.B.;	
"Isolation and structures of alligator gar (Lepisosteus spatula)	
insulin and pancreatic polypeptide.";	
Gen. Comp. Endocrinol. 67:375-382(1987); [2]	
SEQUENCE, AND SYNTHESIS.	
SPECIES=S.canicula; TISSUE=Pancreas;	
MEDLINE=91209266; PubMed=2019251;	
Conlon J.M., Balasubramanian A., Hazen N.;	
"Structural characterization and biological activity of a	
neuropeptide Y-related peptide from the dogfish, Scyliorhinus	
canicula";	
Endocrinology 128:2273-2279(1991). [3]	
SEQUENCE.	
SPECIES=S.acanthias; TISSU=Pancneas;	
Pan J.-Z., Shaw C., Halton D.W., Thim L., Johnston C.F.,	
Fairweather I., Buchanan K.D.;	
"Isolation and primary structure of the peptide Y from the pancreas	
of the spiny dogfish, Squalus acanthias.";	
Regul. Pept. 35:252-252(1991).	
-!- FUNCTION: ELICITS AN INCREASE IN ARTERIAL BLOOD PRESSURE.	
-!- SUBCELLULAR LOCATION: Secretd.	
-!- SIMILARITY: BELONGS TO THE NPY / PPY / PPF FAMILY.	

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DR PIR: S07215; PCGXA.
DR PIR: A60022; PCDFY.
DR PIR: A49743; A49743.
DR HSSP: P01303; IRON.
DR InterPro: IPR001955; Pancreatic_hormn.
DR Pfam: PF00159; hormone3; 1.
DR PRINTS: PR00278; PANCHORMONE.
DR ProDom: PD001267; Pancreatic_hormn; 1.
DR SMART: SM00309; PAH; 1.
DR PROSITE: PS00265; PANCREATIC_HORMONE_1; 1.
DR PROSITE: PS0276; PANCREATIC_HORMONE_2; 1.
KW Hormone: Amidation. 36
FT MOD_RES 36
SQ SEQUENCE 36 AA; 4291 MW; 56A6D8CC086660AA CRC64;

Query Match      80.4%; Score 156; DB 1; Length 36;
Best Local Similarity 75.0%; Pred. No. 5e-15;
Matches 27; Conservative 5; Mismatches 4; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPEELNRYASLRHYLNLTQRY 36
   || ||| ||||| |||| :||:||||:|||||
Db 1 YPPKPNPGEDAPPEELAKYYSALRHYLNLTQRY 36

RESULT 9
PYV_ONCKI
ID PYV_ONCKI STANDARD; PRT; 36 AA.
AC P09474;
DT 01-MAR-1989 (Rel. 10, Created)
DT 01-MAR-1989 (Rel. 10, Last sequence update)
DT 16-OCT-2001 (Rel. 40, Last annotation update)
DE Peptide YY-like (PYY).
OS Oncorhynchus kisutch (Coho salmon), and
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Actinopterygii; Neopterygii; Teleostei; Euteleostei;
OC Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.
OX NCBI_TaxID=8019, 8022;
RN [1]
RP SEQUENCE.
RC SPECIES=O. kisutch; TISSUE=Pancreas;
RX MEDLINE=87128023; PubMed=3545195;
RA Kimmel J.B., Pflsetskaya E.M., Pollock H.G., Hamilton J.W.,
Rouse J.B., Ebner K.E., Rawitch A.B.;
RT "Structure of a peptide from coho salmon endocrine pancreas with
homology to neuropeptide Y.";
RL Biochem. Biophys. Res. Commun. 141:1084-1091(1986).
RN [2]
RP SEQUENCE.
RC SPECIES=O. mykiss; TISSUE=Brain, and Stomach;
RX MEDLINE=93092973; PubMed=1459125;
RA Jensen J., Conlon J.M.;
RT "Characterization of peptides related to neuropeptide tyrosine and
peptide tyrosine-tyrosine from the brain and gastrointestinal tract
of teleost fish.";
RL Eur. J. Biochem. 210:405-410(1992).
RN [3]
RP SEQUENCE.
RC SPECIES=O. mykiss; TISSUE=Brain;
RX MEDLINE=93157164; PubMed=1494498;
RA Barton C.L., Shaw C., Halton D.W., Thim L.;
RT "Rainbow trout (Oncorhynchus mykiss) neuropeptide Y.";
RL Peptides 13:1159-1163(1992).
CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- SIMILARITY: BELONGS TO THE NPY / PYY / PPY FAMILY.
DR PIR: A26377; A26377.
DR HSSP: P01303; IRON.
DR InterPro: IPR001955; Pancreatic_hormn.
DR Pfam: PF00159; hormone3; 1.
DR PRINTS: PR00278; PANCHORMONE.
DR ProDom: PD001267; Pancreatic_hormn; 1.
DR SMART: SM00309; PAH; 1.
DR PROSITE: PS00265; PANCREATIC_HORMONE_1; 1.

DR PIR: S07215; PCGXA.
DR PIR: A60022; PCDFY.
DR PIR: A49743; A49743.
DR HSSP: P01303; IRON.
DR InterPro: IPR001955; Pancreatic_hormn.
DR Pfam: PF00159; hormone3; 1.
DR PRINTS: PR00278; PANCHORMONE.
DR ProDom: PD001267; Pancreatic_hormn; 1.
DR SMART: SM00309; PAH; 1.
DR PROSITE: PS00265; PANCREATIC_HORMONE_1; 1.
DR PROSITE: PS0276; PANCREATIC_HORMONE_2; 1.
KW Hormone: Amidation. 36
FT MOD_RES 36
SQ SEQUENCE 36 AA; 4305 MW; 56A6D8CC08666671 CRC64;

Query Match      79.9%; Score 155; DB 1; Length 36;
Best Local Similarity 75.0%; Pred. No. 6.9e-15;
Matches 27; Conservative 4; Mismatches 5; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPEELNRYASLRHYLNLTQRY 36
   || ||| ||||| |||| :||:||||:|||||
Db 1 YPPKPNPGEDAPPEELAKYYSALRHYLNLTQRY 36

RESULT 10
PYV_RAJRH
ID PYV_RAJRH STANDARD; PRT; 36 AA.
AC P29206;
DT 01-DEC-1992 (Rel. 24, Created)
DT 01-DEC-1992 (Rel. 24, Last sequence update)
DT 16-OCT-2001 (Rel. 40, Last annotation update)
DE Peptide YY-like (PYY).
OS Raja rhina (Skate).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes;
OC Elasmobranchii; Squalia; Hyphosqualea; Pristiogalea; Batoidea;
OC Rajiformes; Rajidae; Raja.
OX NCBI_TaxID=30478;
RN [1]
RP SEQUENCE.
RX MEDLINE=91296574; PubMed=2067973;
RA Conlon J.M., Bjening C., Moon T.W., Youson J.H., Thim L.;
RT "Neuropeptide Y-related peptides from the pancreas of a teleostean
(eel), holostean (bowfin) and elasmobranch (skate) fish.";
RL Peptides 12:221-226(1991).
CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- SIMILARITY: BELONGS TO THE NPY / PYY / PPY FAMILY.
DR HSSP: P01303; IRON.
DR InterPro: IPR001955; Pancreatic_hormn.
DR Pfam: PF00159; hormone3; 1.
DR PRINTS: PR00278; PANCHORMONE.
DR ProDom: PD001267; Pancreatic_hormn; 1.
DR SMART: SM00309; PAH; 1.
DR PROSITE: PS00265; PANCREATIC_HORMONE_1; 1.
DR PROSITE: PS0276; PANCREATIC_HORMONE_2; 1.
KW Hormone: Amidation. 36
FT MOD_RES 36
SQ SEQUENCE 36 AA; 4251 MW; 07A7D9DC196660B6 CRC64;

Query Match      79.9%; Score 155; DB 1; Length 36;
Best Local Similarity 72.2%; Pred. No. 6.9e-15;
Matches 26; Conservative 7; Mismatches 3; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPEELNRYASLRHYLNLTQRY 36
   || ||| ||||| |||| :||:||||:|||||
Db 1 YPPKPNPGDDAPEELAKYYSALRHYLNLTQRY 36

RESULT 11
PYV_BRARE
ID PYV_BRARE STANDARD; PRT; 97 AA.
AC Q918P2;
DT 15-JUN-2002 (Rel. 41, Created)
DT 15-JUN-2002 (Rel. 41, Last sequence update)
DT 15-JUN-2002 (Rel. 41, Last annotation update)
DE Peptide YY precursor.
GN PYY.
OS Brachydanio rerio (Zebrafish) (Danio rerio).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;
OC Cyprinidae; Danio.
OX NCBI_TaxID=7955;
RN [1]
RP SEQUENCE FROM N.A.
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RX MEDLINE=20396163; PubMed=10936170;
RA Soderberg C., Wraith A., Ringvall M., Yan Y.L., Postlethwait J.H.,
RA Brodin L., Larhammar D.;
RT "Zebrafish genes for neuropeptide Y and peptide YY reveal origin by
RT chromosome duplication from an ancestral gene linked to the homeobox
RT cluster.";
RL J. Neurochem. 75:908-918(2000).
CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- SIMILARITY: BELONGS TO THE NPY / PPY / PPY / PPY FAMILY.
CC -----
CC This SWISS-PROT entry is copyright. It is produced through a collaboration
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CC -----
DR EMBL; AF233875; AAF79942.1; -.
DR HSSP; P01303; IRON.
DR ZFIN; ZDB-GENE-980526-71; ppy.
DR InterPro; IPR001955; Pancreatic_hormn.
DR Pfam; PF00159; hormones; 1.
DR PRINTS; PR00278; PANCHORMONE.
DR PRODOM; PD001267; Pancreatic_hormn; 1.
DR SMART; SM00309; PAH; 1.
DR PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.
DR PROSITE; PS00276; PANCREATIC_HORMONE_2; 1.
DR Hormone; Cleavage on pair of basic residues; Amidation; Signal;
KW Neuropeptide.
FT SIGNAL 1 28
FT CHAIN 29 64
FT PROPEP 68 97
FT MOD_RES 64 64
FT SEQUENCE 97 AA; 11175 MW; 96EA07EF0991AC2D CRC64;
Query Match 79.4%; Score 154; DB 1; Length 97;
Best Local Similarity 72.2%; Pred. No. 3,1e-14;
Matches 26; Conservative 6; Mismatches 4; Indels 0; Gaps 0;
QY 1 YPIKPEAGEDASPELNRYASLRHYNLVTRQRY 36
Db 29 YPKPKENPGDAAPELAKYVYALRHYNLVTRQRY 64
RESULT 12
NEUY_DICLA
ID NEUY_DICLA STANDARD; PRT; 99 AA.
AC Q9PTA0; Q9PT97;
DT 16-OCT-2001 (Rel. 40, Created)
DT 16-OCT-2001 (Rel. 40, Last sequence update)
DT 15-JUN-2002 (Rel. 41, Last annotation update)
DE Neuropeptide Y precursor (NPY).
GN NPY.
OS Dicertrarchus labrax (European sea bass).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Actinopterygii; Neopterygii; Teleostei; Euteleostei;
OC Acanthomorpha; Acanthopterygii; Percormorpha; Perciformes; Percoidae;
OC Moronidae; Dicertrarchus.
OX NCBI_TaxID=13489;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=Brain;
RA Cerda-Reverter J.M., Martinez-Rodriguez G., Zanuy S., Carrillo M.,
RA Larhammar D.;
RT "Neuropeptide Y, endocrine gut peptide YY and fish pancreatic peptide
RT Y expression in the brain of a teleost fish (Dicertrarchus labrax):
RT from cloning to evolutionary considerations.";
RL Submitted (APR-1998) to the EMBL/GenBank/DBJ databases.
[2]
RN SEQUENCE OF 1-62 FROM N.A.
RC TISSUE=Blood;

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RA Cerda-Reverter J.M., Martinez-Rodriguez G., Zanuy S., Carrillo M.,
RA Larhammar D.;
RT "Deduced peptide sequence of neuropeptide Y exon 2 from sea bass
RT (Dicertrarchus labrax).";
RL Submitted (APR-1998) to the EMBL/GenBank/DBJ databases.
CC -!- FUNCTION: NPY IS IMPLICATED IN THE CONTROL OF FEEDING AND IN
CC SECRETION OF GONADOTROPHIN-RELEASE HORMONE.
CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- SIMILARITY: BELONGS TO THE NPY / PPY / PPY / PPY FAMILY.
CC -----
CC This SWISS-PROT entry is copyright. It is produced through a collaboration
CC between the Swiss Institute of Bioinformatics and the EMBL outstation -
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CC or send an email to license@isb-sib.ch).
CC -----
DR EMBL; AJ005378; CAB64932.1; -.
DR EMBL; AJ005381; CAB64935.1; -.
DR HSSP; P01303; IRON.
DR InterPro; IPR001955; Pancreatic_hormn.
DR Pfam; PF00159; hormones; 1.
DR PRINTS; PR00278; PANCHORMONE.
DR PRODOM; PD001267; Pancreatic_hormn; 1.
DR SMART; SM00309; PAH; 1.
DR PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.
DR PROSITE; PS00276; PANCREATIC_HORMONE_2; 1.
DR Neuropeptide; Cleavage on pair of basic residues; Signal; Amidation.
FT SIGNAL 1 28
FT CHAIN 29 64
FT PROPEP 68 99
FT MOD_RES 64 64
FT SEQUENCE 99 AA; 11260 MW; 4EEFAED164964184 CRC64;
Query Match 79.4%; Score 154; DB 1; Length 99;
Best Local Similarity 72.2%; Pred. No. 3,2e-14;
Matches 26; Conservative 6; Mismatches 4; Indels 0; Gaps 0;
QY 1 YPIKPEAGEDASPELNRYASLRHYNLVTRQRY 36
Db 29 YPKPKENPGDAAPELAKYVYALRHYNLVTRQRY 64
RESULT 13
NEUY_ONCMY
ID NEUY_ONCMY STANDARD; PRT; 36 AA.
AC P29071;
DT 01-DEC-1992 (Rel. 24, Created)
DT 01-DEC-1992 (Rel. 24, Last sequence update)
DT 15-JUN-2002 (Rel. 41, Last annotation update)
DE Neuropeptide Y (NPY).
GN NPY.
OS Oncorhynchus mykiss (Rainbow trout) (Salmo gairdneri).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Actinopterygii; Neopterygii; Teleostei; Euteleostei;
OC Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.
OX NCBI_TaxID=8022;
RN [1]
RP SEQUENCE.
RC TISSUE=Brain;
RX MEDLINE=93092973; PubMed=1459125;
RA Jensen J., Conlon J.M.;
RT "Characterization of peptides related to neuropeptide tyrosine and
RT peptide tyrosine-tyrosine from the brain and gastrointestinal tract
RT of teleost fish.";
RL Eur. J. Biochem. 210:405-410(1992).
CC -!- FUNCTION: NPY IS IMPLICATED IN THE CONTROL OF FEEDING AND IN
CC SECRETION OF GONADOTROPHIN-RELEASE HORMONE.
CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- SIMILARITY: BELONGS TO THE NPY / PPY / PPY / PPY FAMILY.
DR HSSP; P01303; IRON.
DR InterPro; IPR001955; Pancreatic_hormn.

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DR Pfam: PF00159; hormone3; 1.
DR PRINTS; PR001267; PANCHORMONE.
DR ProDom; PD001267; Panchreatic_hormn; 1.
DR SMART; SM00309; PAH; 1.
DR PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.
DR PROSITE; PS00276; PANCREATIC_HORMONE_2; 1.
DR Neuropeptide; Amidation.
KW MOD_RES 36
SQ SEQUENCE 36 AA; 4311 MW; E2A32293A866611C CRC64;

Query Match      78.9%; Score 153; DB 1; Length 36;
Best Local Similarity 72.2%; Pred. No. 1.3e-14;
Matches 26; Conservative 5; Mismatches 5; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPEELNRYASLRHLYNLVTRQRY 36
   ||||| ||||| ||| :|| :|||:|||||
Db 1 YPVKPNPGEDAPEELAKYKYLALRYHLYNLVTRQRY 36

RESULT 14
PYV_RANRI          STANDARD;          PRT;          36 AA.
AC P29204;
DT 01-DEC-1992 (Rel. 24, Created)
DT 01-DEC-1992 (Rel. 24, Last sequence update)
DT 16-OCT-2001 (Rel. 40, Last annotation update)
DE Peptide YY-like (PYY).
OS Rana ridibunda (Laughing frog) (Marsh frog).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Amphibia; Batrachia; Anura; Neobatrachia; Ranioidea; Ranidae; Rana.
OX NCBI_TaxID=8406;
RN [1]
RP SEQUENCE.
RC TISSUE=Intestine;
RX MEDLINE=92319697; PubMed=1620652;
RA Conlon J.M., Chartrel N., Vaudry H.;
RT "Primary structure of frog PYY: implications for the molecular
   evolution of the pancreatic polypeptide family.";
RL Peptides 13:145-149(1992).
CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- SIMILARITY: BELONGS TO THE NPY / PYY / PYY FAMILY.
DR HSP; P01303; IRON.
DR InterPro; IPR001955; Panchreatic_hormn.
DR Pfam; PF00159; hormone3; 1.
DR PRINTS; PR001267; PANCHORMONE.
DR ProDom; PD001267; Panchreatic_hormn; 1.
DR SMART; SM00309; PAH; 1.
DR PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.
DR PROSITE; PS00276; PANCREATIC_HORMONE_2; 1.
KW Hormone; Amidation.
FT MOD_RES 36
SQ SEQUENCE 36 AA; 4279 MW; 0725C3C6304F663 CRC64;

Query Match      78.9%; Score 153; DB 1; Length 36;
Best Local Similarity 75.0%; Pred. No. 1.3e-14;
Matches 27; Conservative 4; Mismatches 5; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPEELNRYASLRHLYNLVTRQRY 36
   ||||| ||||| ||| :|| :|||:|||||
Db 1 YPPKPNPGEDASPEETKYLALRYHLYNLVTRQRY 36

RESULT 15
NEUY_GADMO          STANDARD;          PRT;          36 AA.
AC P80167;
DT 01-DEC-1992 (Rel. 24, Created)
DT 01-DEC-1992 (Rel. 24, Last sequence update)
DT 15-JUN-2002 (Rel. 41, Last annotation update)
DE Neuropeptide Y (NPY).
GN NPY.
OS Gadus morhua (Atlantic cod).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

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OC Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;
OC Acanthomorpha; Paracanthopterygii; Gadiformes; Gadidae; Gadus.
OX NCBI_TaxID=8049;
RN [1]
RP SEQUENCE.
RC TISSUE=Brain;
RX MEDLINE=93092973; PubMed=1459125;
RA Jensen J., Conlon J.M.;
RT "Characterization of peptides related to neuropeptide tyrosine and
   peptide tyrosine-tyrosine from the brain and gastrointestinal tract
   of teleost fish.";
RL Eur. J. Biochem. 210:405-410(1992).
CC -!- FUNCTION: NPY IS IMPLICATED IN THE CONTROL OF FEEDING AND IN
   SECRETION OF GONADOTROPHIN-RELEASE HORMONE.
CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- SIMILARITY: BELONGS TO THE NPY / PYY / PYY FAMILY.
DR PIR; S27054; S27054.
DR HSP; P01303; IRON.
DR InterPro; IPR001955; Panchreatic_hormn.
DR Pfam; PF00159; hormone3; 1.
DR PRINTS; PR001267; PANCHORMONE.
DR ProDom; PD001267; Panchreatic_hormn; 1.
DR SMART; SM00309; PAH; 1.
DR PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.
DR PROSITE; PS00276; PANCREATIC_HORMONE_2; 1.
KW Neuropeptide; Amidation.
FT MOD_RES 36
SQ SEQUENCE 36 AA; 4267 MW; 17B09AA83867A7B6 CRC64;

Query Match      78.4%; Score 152; DB 1; Length 36;
Best Local Similarity 72.2%; Pred. No. 1.8e-14;
Matches 26; Conservative 6; Mismatches 4; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPEELNRYASLRHLYNLVTRQRY 36
   ||||| ||||| ||| :|| :|||:|||||
Db 1 YPIKPNPGEDAPADELAKYYSALRHYINLVTRQRY 36

Search completed: February 27, 2003, 14:36:51
Job time : 30 secs

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Result No.	Score	Query ⁸		Length	DB	ID	Description
		Match					
1	188	96.9	36	6	Q9TR93	Q9tr93 oryctolagus	
2	184	94.8	98	11	Q91XD0	Q91xd0 mus musculus	
3	176	90.7	34	6	Q9TR92	Q9tr92 oryctolagus	
4	151	77.8	99	13	Q90WF4	Q90wf4 paralichthy	
5	149	76.8	95	13	Q919D3	Q919d3 ictalurus p	
6	147	75.8	96	13	Q9DGK7	Q9dgk7 cyprinus ca	
7	146	75.3	76	6	Q9N0M5	Q9nm05 sus scrofa	
8	144	74.2	97	6	Q9XSW6	Q9xsw6 macaca mula	
9	143	73.7	89	11	Q92SV2	Q92sv2 mus musculus	
10	143	73.7	90	6	Q9TS16	Q9ts16 ovis aries	
11	143	73.7	97	6	Q8SPR7	Q8spr7 ovis aries	
12	141	72.7	97	13	Q9PW68	Q9pw68 typhlonecte	
13	141	72.7	99	13	Q90WF3	Q90wf3 paralichthy	
14	136	70.1	36	13	Q9PS46	Q9ps46 scyllorhinu	
15	132	68.0	97	13	Q90WF2	Q90wf2 paralichthy	
16	98	50.5	59	6	Q9GK10	Q9gk10 sus scrofa	

Db 2 KPEAPGSDASPEELNRYYSALRHVNLVTRQY 34

RESULT 4

Q90WF4 PRELIMINARY; PRT; 99 AA.

ID Q90WF4

AC Q90WF4; 19, Created)

DT 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)

DT 01-MAR-2002 (TrEMBLrel. 20, Last annotation update)

DE Neuropeptide Y.

GN NPY.

OS Paralicthys olivaceus (Flounder).

OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

OC Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;

OC Acanthomorpha; Acanthopterygii; Percomorpha; Pleuronectiformes;

OC Pleuronectoidae; Paralicthiidae; Paralicthys.

OX NCBI_TaxID=8255;

OS NCBI_TaxID=8255;

RN [1]

RP SEQUENCE FROM N.A.

RA Kurokawa T., Suzuki T.

RT "Development of neuropeptide Y related peptides in the digestive

RT organs during the larval stage of Japanese flounder, Paralicthys

RT olivaceus."

RL Submitted (FEB-2001) to the EMBL/GenBank/DBJ databases.

DR EMBL: AB055211; BAB62409.1; -.

DR InterPro: IPR001955; Pancreatc_hormn.

DR Pfam: PF00159; hormone3; 1.

DR ProDom: PD001267; Pancreatc_hormn; 1.

DR PROSITE; PS00265; PANCREATIC_HORMONE_1; UNKNOWN_1.

DR PROSITE; PS0276; PANCREATIC_HORMONE_2; 1.

SQ SEQUENCE 99 AA; 11215 MW, 6FEED47F24CF6498 CRC64;

Query Match 77.8%; Score 151; DB 13; length 99;

Best Local Similarity 69.4%; Pred. No. 7.1e-13;

Matches 25; Conservative 7; Mismatches 4; Indels 0; Gap

QY 1 YPKPKEAPGSDASPEELNRYYSALRHVNLVTRQY 36

DB 29 YPKPKEAPGSDASPEELNRYYSALRHVNLVTRQY 64

QY 1 YPKPKEAPGSDASPEELNRYYSALRHVNLVTRQY 36

DB 29 YPKPKEAPGSDASPEELNRYYSALRHVNLVTRQY 64

RESULT 5

Q919D3 PRELIMINARY; PRT; 95 AA.

ID Q919D3

AC Q919D3; 15, Created)

DT 01-OCT-2000 (TrEMBLrel. 15, Last sequence update)

DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)

DE Neuropeptide Y.

OS Ictalurus punctatus (Channel catfish).

OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

OC Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Siluriformes;

OC Ictaluridae; Ictalurus.

OX NCBI_TaxID=7998;

RN [1]

RP SEQUENCE FROM N.A.

RA Leonard J.B.K., Waldbieser G.C., Silverstein J.T.;

RT "Neuropeptide Y (NPY) sequence and distribution in channel catfish

RT (Ictalurus punctatus).";

RL Submitted (MAY-2000) to the EMBL/GenBank/DBJ databases.

CC -1- SIMILARITY: BELONGS TO THE NPY / PPY / PYY FAMILY.

DR EMBL: AF267164; AAF1617.1; -.

DR HSSP; P01303; IRON.

DR InterPro: IPR001955; Pancreatc_hormn.

DR Pfam: PF00159; hormone3; 1.

DR PRINTS; PR00278; PANCHORMONE.

DR ProDom: PD001267; Pancreatc_hormn; 1.

DR SMART; SM00309; PAH; 1.

DR PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.

DR PROSITE; PS0276; PANCREATIC_HORMONE_2; 1.

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RESULT 2
Q91XD0 Q91XD0 PRELIMINARY; PRT; 98 AA.
ID AC Q91XD0;
DT 01-DEC-2001 (TREMBlrel. 19, Created)
DT 01-DEC-2001 (TREMBlrel. 19, Last sequence update)
DT 01-MAR-2002 (TREMBlrel. 20, Last annotation update)
DE Unknown (protein for MGC:19143).
OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OC NCBI_TaxID=10090;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE-COLON;
RA Strausberg R.;
RL Submitted (JUL-2001) to the EMBL/GenBank/DBDJ databases.
DR EMBL; BC010821; AAH10821.1; -.
DR InterPro; IPR001955; Pancreatc_hormn.
DR Pfam; PF00159; hormone3; 1.
DR ProDom; PD001267; Pancreatc_hormn; 1.
DR PROSITE; PS00265; PANCREATIC_HORMONE_1; UNKNOWN_1.
DR PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.
SQ SEQUENCE 98 AA; 11064 MW; 7AF165A1052C3249 CRC64;

Query Match 94.98; Score 184; DB 11; Length 98;
Best Local Similarity 94.48; Pred. No. 2.4e-17;
Matches 34; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 1 YPIKPEAGDASPEELNRYVYASLRHYLNLVTRQRY 36
|| |||||
Db 29 YPARPEAGDASPEELSRYYASLRHYLNLVTRQRY 64

RESULT 3
Q9TR92 Q9TR92 PRELIMINARY; PRT; 34 AA.
ID AC Q9TR92;
DT 01-MAY-2000 (TREMBlrel. 13, Created)
DT 01-MAY-2000 (TREMBlrel. 13, Last sequence update)
DT 01-OCT-2001 (TREMBlrel. 16, Last annotation update)
DE Peptide YY, PYY(3-36).
OS Oryctolagus cuniculus (Rabbit).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
OC NCBI_TaxID=9986;
RN [1]
RP SEQUENCE.
RX MEDLINE=95075735; PubMed=7984499;
RA Grandt D., Schmiczek M., Struk K., Shively J., Eysselein V.E.,
RA Goebell H., Reeve J.R.Jr.;
RT "Characterization of two forms of peptide YY, PYY(1-36) and PYY(3-36),
RT in the rabbit."
RL Peptides 15:815-820(1994).
CC -1- SIMILARITY: BELONGS TO THE NPY / PYY / PYY FAMILY.
DR HSSP; P01303; IRON.
DR InterPro; IPR001955; Pancreatc_hormn.
DR Pfam; PF00159; hormone3; 1.
DR ProDom; PD001267; Pancreatc_hormn; 1.
DR SMART; SM00309; PAH; 1.
DR PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.
DR PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.
KW Animation.
SQ SEQUENCE 34 AA; 4024 MW; 02D4E9C38BA5FC8D CRC64;

Query Match 90.78; Score 176; DB 6; Length 34;
Best Local Similarity 100.0%; Pred. No. 9.1e-17;
Matches 33; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 4 KPEAPGEDASPEELNRYVYASLRHYLNLVTRQRY 36
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[illegible]

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OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX NCBI_TaxID=10090;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=SM/J;
RX MEDLINE=21077529; PubMed=11210195;
RA Taylor B.A., Wnek C., Schroeder D., Phillips S.J.;
RT "Multiple obesity Otis identified in an intercross between the NZO
RT (New Zealand obese) and the SM (small) mouse strains.";
RL Mamm. Genome 12:95-103(2001).
DR EMBL; AF286198; AAG01330.1; -
DR InterPro: IPR001955; Pancreatic_hormn.
DR Pfam: PF00159; hormone3; 1.
DR ProDom; PD001267; Pancreatic_hormn; 1.
DR PROSITE; PS00265; PANCREATIC_HORMONE_1; UNKNOWN_1.
DR PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.
FT NON_TER 89
SQ SEQUENCE 89 AA; 9943 MW; AE6052615A59D96A CRC64;

Query Match 73.7%; Score 143; DB 11; Length 89;
Best Local Similarity 66.7%; Pred. No. 7.7e-12;
Matches 24; Conservative 7; Mismatches 5; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPEELNRYVYASLRHYLNLVTRQRY 36
   11 11: 11111 1: 11111111111111111111
Db 29 YPSKPNPGDAPEDMRYVYASLRHYLNLVTRQRY 64

RESULT 10
Q9TS16 PRELIMINARY; PRT; 90 AA.
AC Q9TS16;
DT 01-MAY-2000 (TrEMBLrel. 13, Created)
DT 01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
DE Neuropeptide Y (Fragment).
OS Ovis aries (Sheep).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae;
OC Bovidae; Caprinae; Ovis.
OX NCBI_TaxID=9940;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=SM/J;
RX MEDLINE=21077529; PubMed=11210195;
RA Taylor B.A., Wnek C., Schroeder D., Phillips S.J.;
RT "Multiple obesity Otis identified in an intercross between the NZO
RT (New Zealand obese) and the SM (small) mouse strains.";
RL Mamm. Genome 12:95-103(2001).
DR EMBL; AF286198; AAG01330.1; -
DR InterPro: IPR001955; Pancreatic_hormn.
DR Pfam: PF00159; hormone3; 1.
DR ProDom; PD001267; Pancreatic_hormn; 1.
DR PROSITE; PS00265; PANCREATIC_HORMONE_1; UNKNOWN_1.
DR PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.
FT NON_TER 89
SQ SEQUENCE 89 AA; 9943 MW; AE6052615A59D96A CRC64;

Query Match 73.7%; Score 143; DB 11; Length 89;
Best Local Similarity 66.7%; Pred. No. 7.7e-12;
Matches 24; Conservative 7; Mismatches 5; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPEELNRYVYASLRHYLNLVTRQRY 36
   11 11: 11111 1: 11111111111111111111
Db 29 YPSKPNPGDAPEDMRYVYASLRHYLNLVTRQRY 64

RESULT 10
Q9TS16 PRELIMINARY; PRT; 90 AA.
AC Q9TS16;
DT 01-MAY-2000 (TrEMBLrel. 13, Created)
DT 01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
DE Neuropeptide Y (Fragment).
OS Ovis aries (Sheep).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae;
OC Bovidae; Caprinae; Ovis.
OX NCBI_TaxID=9940;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=SM/J;
RX MEDLINE=21077529; PubMed=11210195;
RA Taylor B.A., Wnek C., Schroeder D., Phillips S.J.;
RT "Multiple obesity Otis identified in an intercross between the NZO
RT (New Zealand obese) and the SM (small) mouse strains.";
RL Mamm. Genome 12:95-103(2001).
DR EMBL; AF286198; AAG01330.1; -
DR InterPro: IPR001955; Pancreatic_hormn.
DR Pfam: PF00159; hormone3; 1.
DR ProDom; PD001267; Pancreatic_hormn; 1.
DR PROSITE; PS00265; PANCREATIC_HORMONE_1; UNKNOWN_1.
DR PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.
FT NON_TER 89
SQ SEQUENCE 89 AA; 9943 MW; AE6052615A59D96A CRC64;

Query Match 73.7%; Score 143; DB 6; Length 97;
Best Local Similarity 66.7%; Pred. No. 8.4e-12;
Matches 24; Conservative 7; Mismatches 5; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPEELNRYVYASLRHYLNLVTRQRY 36
   11 11: 11111 1: 11111111111111111111
Db 29 YPSKPNPGDAPEDMRYVYASLRHYLNLVTRQRY 64

RESULT 12
Q9PW68 PRELIMINARY; PRT; 97 AA.
AC Q9PW68;
DT 01-MAY-2000 (TrEMBLrel. 13, Created)
DT 01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
DT 01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
DE Neuropeptide Y precursor (NPY).
OS NPY.
OC Typhlonectes natans (Rio Cauca caecilian).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Amphibia; Gymnophiona; Caeciliidae; Typhlonectes.
OX NCBI_TaxID=8456;
RN [1]
RP SEQUENCE FROM N.A.; SEQUENCE OF 29-64, TISSUE SPECIFICITY, AND MASS
RP SPECTROMETRY.
RC TISSUE=BRAIN;
RX MEDLINE=21184972; PubMed=11287086;
RA Ebersole T.J., Conlon J.M., Goetz F.W., Boyd S.K.;
RT "Characterization and distribution of neuropeptide Y in the brain of a
RT caecilian amphibian.";
RL Peptides 22:325-334(2001).
CC -!- FUNCTION: NPY IS IMPLICATED IN THE CONTROL OF FEEDING AND IN
CC SECRETION OF GONADOTROPHIN-RELEASE HORMONE (BY SIMILARITY).
CC -!- SUBCELLULAR LOCATION: SECRETED (BY SIMILARITY).
CC -!- TISSUE SPECIFICITY: EXPRESSED THROUGHOUT THE BRAIN WITH HIGHEST
CC LEVELS OF EXPRESSION IN MEDIAL PALLIUM, BASAL FOREBRAIN, PREOPTIC
CC AREA, MIDBRAIN TEGMENTUM AND TRIGEMINAL NUCLEUS.
CC -!- MASS SPECTROMETRY: MW=4246; MW_ERR=4; METHOD=NALDI.
CC -!- SIMILARITY: BELONGS TO THE NPY / PPY / PPY FAMILY.
DR EMBL; AF167559; AAD48033.1; -
DR HSSP; P01303; IRON.
DR InterPro: IPR001955; Pancreatic_hormn.
DR Pfam: PF00159; hormone3; 1.
DR PRINTS; PR00278; PANCHORMONE.
DR ProDom; PD001267; Pancreatic_hormn; 1.
DR SMART; SM00309; PAH; 1.
DR PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.
DR PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.
KW Amidation.
FT NON_TER 90
FT NON_TER 90
SQ SEQUENCE 90 AA; 9916 MW; 46FF2FB153EE5FFB CRC64;

Query Match 73.7%; Score 143; DB 6; Length 90;
Best Local Similarity 66.7%; Pred. No. 7.8e-12;
Matches 24; Conservative 7; Mismatches 5; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPEELNRYVYASLRHYLNLVTRQRY 36
   11 11: 11111 1: 11111111111111111111
Db 24 YPSKPNPGDAPEDMRYVYASLRHYLNLVTRQRY 59

RESULT 11
Q8SPF7

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ID Q8SPF7 PRELIMINARY; PRT; 97 AA.
AC Q8SPF7;
DT 01-JUN-2002 (TrEMBLrel. 21, Created)
DT 01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
DT 01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
DE Neuropeptide Y precursor.
GN NPY.
OS Ovis aries (Sheep).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae;
OC Bovidae; Caprinae; Ovis.
OX NCBI_TaxID=9940;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=ILE DE FRANCE; TISSUE=HYPOTHALAMUS;
RA Pillon D., Bruneau G.;
RT "Nucleotide sequence of Ovine preproneuropeptide Y.";
RT Submitted (OCT-2001) to the EMBL/GenBank/DBJ databases.
DR EMBL; AJ417904; CAD10677.1; -
DR Signal.
KW SIGNAL.
FT SIGNAL.
FT CHAIN 29
FT CHAIN 68
FT CHAIN 97
SQ SEQUENCE 97 AA; 10750 MW; 6C2209A361CF8583 CRC64;

Query Match 73.7%; Score 143; DB 6; Length 97;
Best Local Similarity 66.7%; Pred. No. 8.4e-12;
Matches 24; Conservative 7; Mismatches 5; Indels 0; Gaps 0;

QY 1 YPIKPEAPGEDASPEELNRYVYASLRHYLNLVTRQRY 36
   11 11: 11111 1: 11111111111111111111
Db 29 YPSKPNPGDAPEDMRYVYASLRHYLNLVTRQRY 64

RESULT 12
Q9PW68 PRELIMINARY; PRT; 97 AA.
AC Q9PW68;
DT 01-MAY-2000 (TrEMBLrel. 13, Created)
DT 01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
DT 01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
DE Neuropeptide Y precursor (NPY).
OS NPY.
OC Typhlonectes natans (Rio Cauca caecilian).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Amphibia; Gymnophiona; Caeciliidae; Typhlonectes.
OX NCBI_TaxID=8456;
RN [1]
RP SEQUENCE FROM N.A.; SEQUENCE OF 29-64, TISSUE SPECIFICITY, AND MASS
RP SPECTROMETRY.
RC TISSUE=BRAIN;
RX MEDLINE=21184972; PubMed=11287086;
RA Ebersole T.J., Conlon J.M., Goetz F.W., Boyd S.K.;
RT "Characterization and distribution of neuropeptide Y in the brain of a
RT caecilian amphibian.";
RL Peptides 22:325-334(2001).
CC -!- FUNCTION: NPY IS IMPLICATED IN THE CONTROL OF FEEDING AND IN
CC SECRETION OF GONADOTROPHIN-RELEASE HORMONE (BY SIMILARITY).
CC -!- SUBCELLULAR LOCATION: SECRETED (BY SIMILARITY).
CC -!- TISSUE SPECIFICITY: EXPRESSED THROUGHOUT THE BRAIN WITH HIGHEST
CC LEVELS OF EXPRESSION IN MEDIAL PALLIUM, BASAL FOREBRAIN, PREOPTIC
CC AREA, MIDBRAIN TEGMENTUM AND TRIGEMINAL NUCLEUS.
CC -!- MASS SPECTROMETRY: MW=4246; MW_ERR=4; METHOD=NALDI.
CC -!- SIMILARITY: BELONGS TO THE NPY / PPY / PPY FAMILY.
DR EMBL; AF167559; AAD48033.1; -
DR HSSP; P01303; IRON.
DR InterPro: IPR001955; Pancreatic_hormn.
DR Pfam: PF00159; hormone3; 1.
DR PRINTS; PR00278; PANCHORMONE.
DR ProDom; PD001267; Pancreatic_hormn; 1.
DR SMART; SM00309; PAH; 1.
DR PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.
DR PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.

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Neuropeptide; Cleavage on pair of basic residues; Signal; Amidation.

	SIGNAL	1	28	
FT	PEPTIDE	29	64	NEUROPEPTIDE Y.
FT	PROPEP	68	97	BY SIMILARITY.
FT	MOD_RES	64		AMIDATION (G-65 PROVIDE AMIDE GROUP)
FT				(BY SIMILARITY).
SEQ	SEQUENCE	97 AA; 11275 MW;	9682E9D89DFB1C7 CRC64;	

Query Match
Best Local Similarity 72.7%; Score 141; DB 13; Length 97;
Matches 23; Conservative 8; Mismatches 5; Indels 0; Gaps 0;

	YPIRPEARAGEDASPEELNRYVYASLRHLYNLVTRQRY	36
Qy	: : : :	
Dd	29 YPSKPDNFGEDAPAEADMAKYYSALRHYNLITRQY	64

RESULT 13

	PRELIMINARY;	PRT;	99 AA.
ID	Q90WF3		
ID	Q90WF3		
DT	01-DEC-2001 (TrEMBLrel. 19, Created)		
DT	01-DEC-2001 (TrEMBLrel. 19, Last sequence update)		
DT	01-MAR-2002 (TrEMBLrel. 20, Last annotation update)		
DE	Peptide YY.		
GN	PYY.		
OS	Paralichthys olivaceus (Flounder).		
OC	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;		
OC	Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;		
OC	Acanthomorpha; Acanthopterygii; Perciformes; Pleuronectiformes;		
OC	Pleuronectoidei; Paralichthyidae; Paralichthys.		
OX	NCBI_TaxID=8255;		
RN	[1]		
RP	SEQUENCE FROM N.A.		
RC	TISSUP-BRAIN;		
RA	Kurokawa T.; Suzuki T.;		
RT	"Development of neuropeptide Y related peptides in the digestive		
RT	organs during the larval stage of Japanese flounder, Paralichthys		
RT	olivaceus."		
RL	Submitted (FEB-2001) to the EMBL/GenBank/DDBJ databases.		
DR	EMBL; AB055212; BAB62410.1; "		
DR	InterPro; IPR001955; Pancreatic_hormn.		
DR	Pfam; PF00159; hormones; 1.		
DR	ProDom; PD001267; Pancreatic_hormn; 1.		
DR	PROSITE; PS00265; PANCREATIC_HORMONE_1; UNKNOWN_1.		
DR	PROSITE; PS00276; PANCREATIC_HORMONE_2; 1		
SEQ	SEQUENCE 99 AA; 11179 MW; 32F6C21217CB1984 CRC64;		

Query Match
Best Local Similarity 72.7%; Score 141; DB 13; Length 99;
Matches 23; Conservative 8; Mismatches 5; Indels 0; Gaps 0;

	YPIRPEARAGEDASPEELNRYVYASLRHLYNLVTRQRY	36
Qy	: : : :	
Dd	28 YPVAKTTPREGATPEDLAKYYSALRHYNLITRQY	63

RESULT 14

	PRELIMINARY;	PRT;	36 AA.
ID	Q9PS46		
ID	Q9PS46		
DT	01-MAY-2000 (TrEMBLrel. 13, Created)		
DT	01-MAY-2000 (TrEMBLrel. 13, Last sequence update)		
DT	01-OCT-2001 (TrEMBLrel. 18, Last annotation update)		
DE	Neuropeptide Y, NP-PANCREATIC polypeptide homolog.		
OS	Scyliorhinus canicula (Spotted dogfish) (Spotted catshark).		
OC	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes;		
OC	Elasmobranchii; Galeomorphii; Galeoidea; Carchariniiformes;		
OC	Scyliorhinidae; Scyliorhinus.		
OX	NCBI_TaxID=7830;		
RN	[1]		
RP	SEQUENCE.		
RX	MEDLINE=92396601; PubMed=1523163;		

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